

“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, 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    Alive    443
## 4 Yes    Dead     139
```

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    Dead      139 0.239
```

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
```

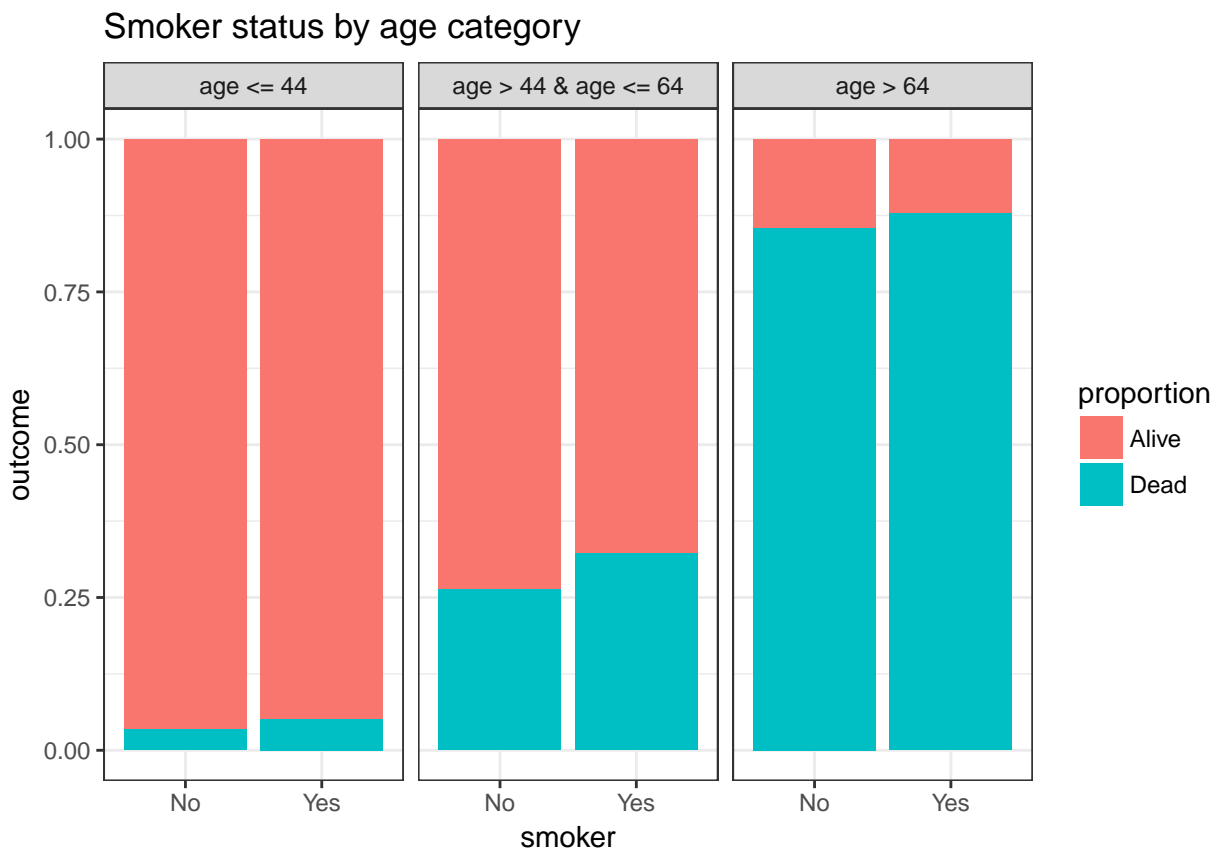
```
##   outcome smoker age      category
## 1  Alive   Yes  23   age <= 44
## 2  Alive   Yes  18   age <= 44
## 3  Dead   Yes  71   age > 64
## 4  Alive   No  67   age > 64
## 5  Alive   No  64 age > 44 & age <= 64
## 6  Alive   Yes  38   age <= 44
```

```
# 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]
##   category      smoker outcome      n proportion
##   <fct>         <fct>   <fct>   <int>     <dbl>
## 1 age <= 44     No     Alive     327     0.965
## 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     Alive     147     0.735
## 6 age > 44 & age <= 64 No     Dead       53     0.265
```

```
# create the visualization with the above result set and facet on age categories
Whickham_cat %>%
ggplot() +
  geom_bar(aes(x = smoker, y = proportion, fill = outcome, label = round(proportion,2)),
           stat = 'identity', position = 'fill') +
  labs(x = 'smoker', y = 'outcome', fill = 'proportion',
       title = 'Smoker status by age category') +
  facet_grid(~ category) +
  theme_bw()
```



Answer:

From the above dataset it is observed that until 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.

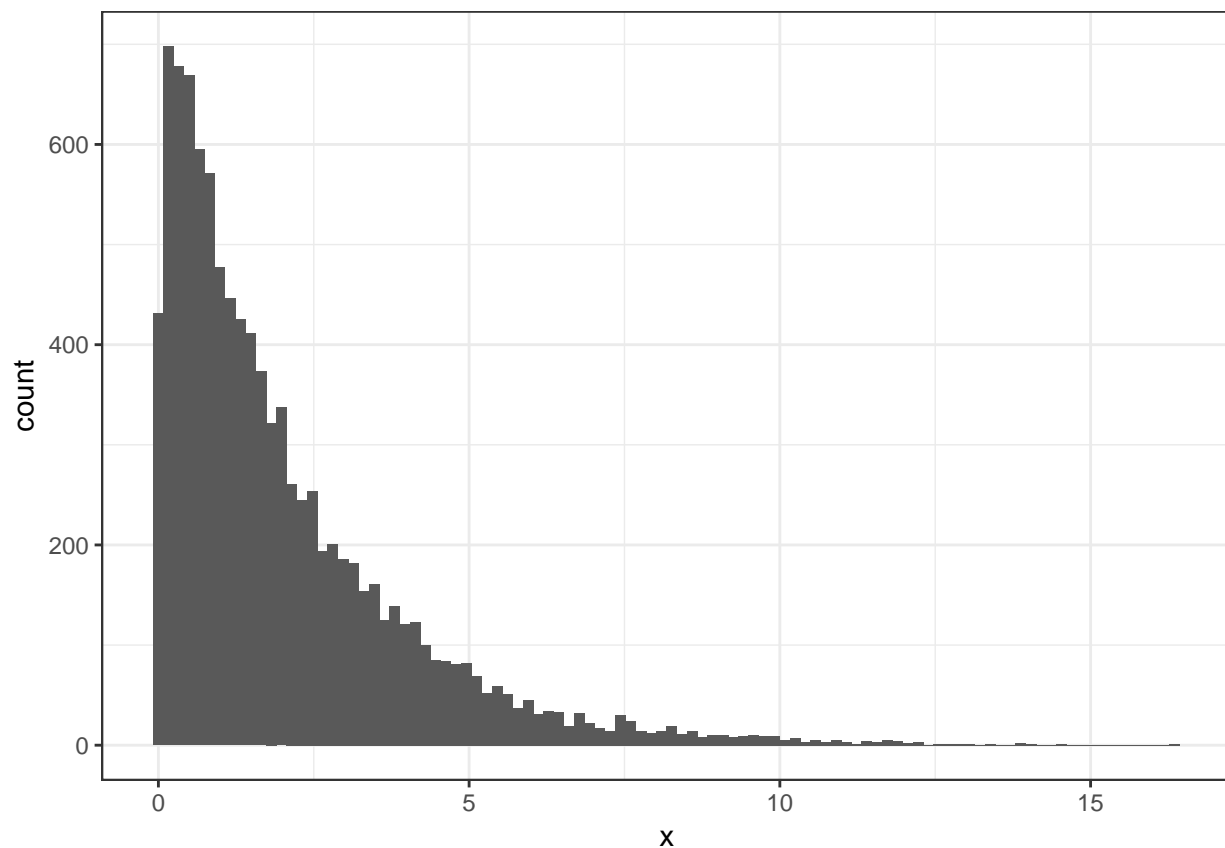
Exercise 2

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
```

```
## # A tibble: 10,000 x 1
##       x
##   <dbl>
## 1 2.70
## 2 0.346
## 3 0.324
## 4 0.00383
## 5 2.67
## 6 0.399
## 7 0.540
## 8 0.262
## 9 1.99
## 10 0.610
## # ... with 9,990 more rows
```

```
# histogram for the above sample gamma
ggplot(data = gamma_samp) +
  geom_histogram(aes(x=x), bins=100) +
  theme_bw()
```



Question: 2

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

```
## [1] "Mean = 2.010743"  
# finding standard deviation  
sprintf("Standard deviation = %f", sapply(gamma_samp, sd, na.rm = TRUE))  
  
## [1] "Standard deviation = 1.981121"
```

Question: 3

Question: 4

Question: 5

Question: 6

Question: 7

Question: 8