



## Exercises: Data Visualisation

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# Chapter 1

# Introduction

In this document you will find some exercises about data visualisation. Most of the exercises are composed by some basic data visualisation questions and some questions on advanced topics of data visulisation, labeled as advanced.

## Chapter 2

# Data Visualization with ggplot2

Load ggplot2 package, supposing it is already installed.

```
require(tidyverse)
require(grid)
```

### 2.1 Data

### 2.1.1 iris

Some of the following exercises are based on the iris dataset, taken from the datasets package. It is a base package so it is already installed and loaded.

```
data("iris")
```

This dataset gives the measurements in centimeters of length and width of sepal and petal, respectively, for 50 flowers from each of 3 species of iris. The species are Iris setosa, versicolor, and virginica.

iris dataset contains the following variables:

```
• Sepal.Length: length of iris sepal
```

• Sepal.Width: width of iris sepal

• Petal.Length: length of iris petal

• Petal.Width: width of iris petal

• Species: species of iris

dim(iris)

```
## [1] 150 5
head(iris)
     Sepal.Length Sepal.Width Petal.Length Petal.Width Species
## 1
              5.1
                          3.5
                                      1.4
                                                  0.2 setosa
## 2
              4.9
                         3.0
                                      1.4
                                                  0.2 setosa
## 3
              4.7
                         3.2
                                      1.3
                                                  0.2 setosa
## 4
              4.6
                         3.1
                                      1.5
                                                  0.2 setosa
## 5
              5.0
                         3.6
                                      1.4
                                                  0.2 setosa
                                                  0.4 setosa
## 6
              5.4
                         3.9
                                      1.7
str(iris)
## 'data.frame':
                    150 obs. of 5 variables:
## $ Sepal.Length: num 5.1 4.9 4.7 4.6 5 5.4 4.6 5 4.4 4.9 ...
## $ Sepal.Width : num 3.5 3 3.2 3.1 3.6 3.9 3.4 3.4 2.9 3.1 ...
## $ Petal.Length: num 1.4 1.4 1.3 1.5 1.4 1.7 1.4 1.5 1.4 1.5 ...
## $ Petal.Width : num 0.2 0.2 0.2 0.2 0.2 0.4 0.3 0.2 0.2 0.1 ...
## $ Species : Factor w/ 3 levels "setosa", "versicolor", ...: 1 1 1 1 1 1 1 1 1 1 ...
```

### 2.1.2 Comic characters data

Other exercises are based on marvel\_wikia\_data dataset, that you may find in the folder exercises/data.

```
marvel_wikia_data <- read_csv("marvel-wikia-data.csv")</pre>
## Parsed with column specification:
## cols(
##
     page_id = col_integer(),
##
     name = col_character(),
##
     urlslug = col_character(),
     ID = col_character(),
##
##
     ALIGN = col_character(),
##
     EYE = col_character(),
##
     HAIR = col_character(),
##
     SEX = col_character(),
##
     GSM = col_character(),
##
     ALIVE = col_character(),
     APPEARANCES = col_integer(),
##
     `FIRST APPEARANCE` = col_character(),
##
     Year = col_integer()
## )
```

2.1. DATA 9

The data comes from Marvel Wikia. The file was scraped in August 2014 and contains the following variables:

- page\_id: The unique identifier for that characters page within the wikia
- name: The name of the character
- urlslug: The unique url within the wikia that takes you to the character
- ID: The identity status of the character (Secret Identity, Public identity, [on marvel only: No Dual Identity])
- ALIGN: If the character is Good, Bad or Neutral
- EYE: Eye color of the character
- HAIR: Hair color of the character
- SEX: Sex of the character (e.g. Male, Female, etc.)
- GSM: If the character is a gender or sexual minority (e.g. Homosexual characters, bisexual characters)
- ALIVE: If the character is alive or deceased
- APPEARANCES: The number of appareances of the character in comic books (as of Sep. 2, 2014. Number will become increasingly out of date as time goes on.)
- FIRST APPEARANCE The month and year of the character's first appearance in a comic book, if available
- YEAR: The year of the character's first appearance in a comic book, if available

```
dim(marvel_wikia_data)
## [1] 16376 13
```

#### head(marvel wikia data)

```
## # A tibble: 6 x 13
##
     page_id
                                                   name
##
       <int>
                                                   <chr>
## 1
        1678
                              Spider-Man (Peter Parker)
## 2
                       Captain America (Steven Rogers)
        7139
## 3
       64786 "Wolverine (James \\\"Logan\\\" Howlett)"
## 4
               "Iron Man (Anthony \\\"Tony\\\" Stark)"
        1868
## 5
        2460
                                    Thor (Thor Odinson)
        2458
                             Benjamin Grimm (Earth-616)
## # ... with 11 more variables: urlslug <chr>, ID <chr>, ALIGN <chr>,
       EYE <chr>, HAIR <chr>, SEX <chr>, GSM <chr>, ALIVE <chr>,
## #
## #
       APPEARANCES <int>, `FIRST APPEARANCE` <chr>, Year <int>
```

## 2.2 Scatterplot

Let us consider iris dataset.

### **2.2.1** Exercise 1

- a. Generate a scatterplot to analyze the relationship between Sepal.Width and Sepal.Length variables.
- b. Set the size of the point as 3 and their colour (colour and fill arguments) as "orchid3".

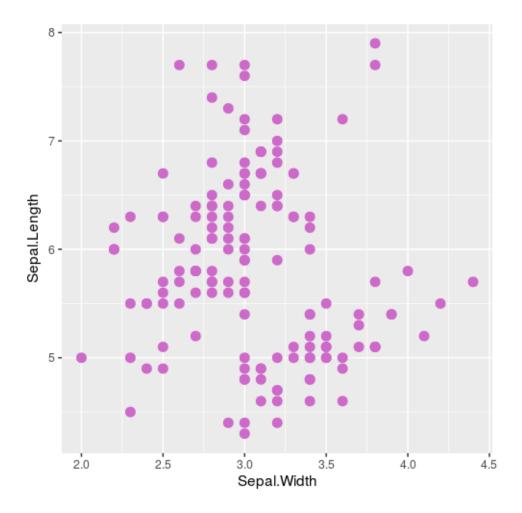


Figure 2.1:

2.2. SCATTERPLOT 11

c. advanced: Add "Sepal Characteristics" as a red italic title and change axis title to "Sepal length" and "Sepal width".

```
pl + ggtitle("Sepal Characteristics") +
  labs(x = "Sepal width", y = "Sepal length") +
  theme(plot.title=element_text(face="italic", colour="red"))
```

## Sepal Characteristics

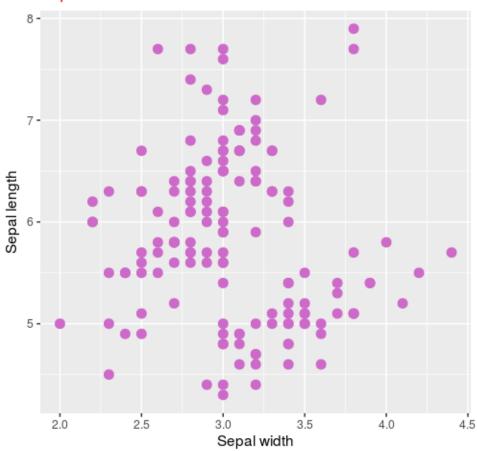


Figure 2.2:

### 2.2.2 Exercise 2

a. Generate a scatterplot to analyze the relationship between Petal.Width and Petal.Length variables according to iris species, mapped as colour aes.

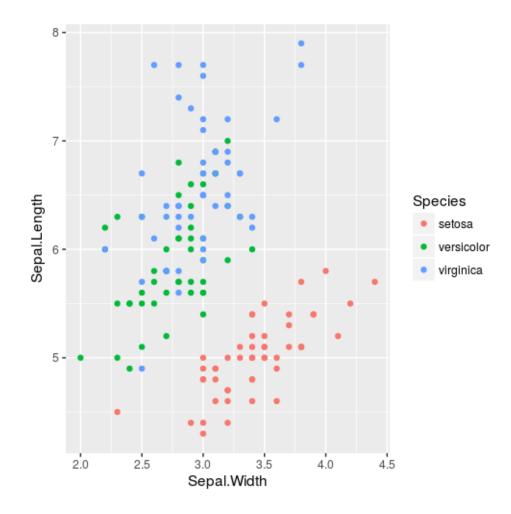


Figure 2.3:

- b. advanced: Change axis title to "Sepal length" and "Sepal width".
- c. advanced: Move the legend to the bottom.

```
pl +
  labs(x = "Sepal width", y = "Sepal length") +
  theme(legend.position="bottom")
```

2.2. SCATTERPLOT 13

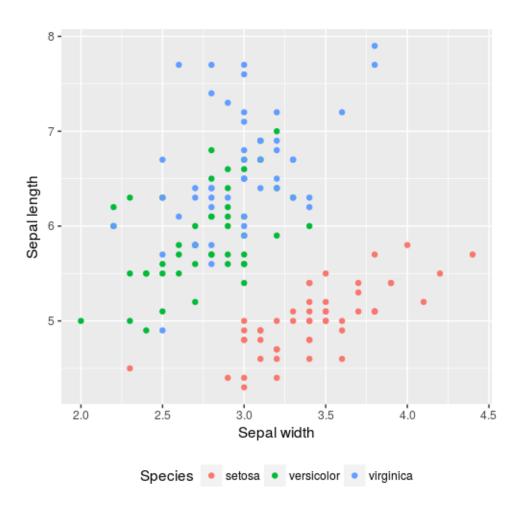


Figure 2.4:

### 2.3 Line PLot

Let us consider marvel\_wikia\_data dataset.

### 2.3.1 Exercise 1

- a. Build a line plot to see the number of new characters that come out each year.
- b. Build a lineplot to compare the differences in the number of female characters and male characters that come out each year.

```
number_characters <- marvel_wikia_data %>%
 group_by(Year, SEX) %>%
 summarise(new_char = n()) %>%
 ungroup()
ggplot(data=number_characters, mapping=aes(x=Year, y=new_char, colour= SEX)) +
 geom_line()
c. Do as in (b.) but use different line types as well as different point types and different
  colours
ggplot(data=number_characters, mapping=aes(x=Year, y=new_char, colour= SEX)) +
 geom_line(mapping=aes(linetype = SEX)) +
 geom_point(mapping=aes(shape = SEX))
d. advanced: Choose a blue colour palette to represent the different lines (use the command
  scale_colour_brewer( palette = "PuBu" ))
e. advanced: Modify axis names and the key labels with scale_colour_brewer choosing
  options name = "Characters gender" and labels = c("Agender", "Female",
   "Genderfluid", "Male", "Not available")).
ggplot(data=number_characters, mapping=aes(x=Year, y=new_char, colour= SEX)) +
 geom_line(mapping=aes(colour = SEX)) + labs(x = "Year", y = "Number of characters") +
 scale_colour_brewer(palette="PuBu", name = "Characters gender",
                      labels=c("Agender", "Female", "Genderfluid",
```

"Male", "Not available"))

2.3. LINE PLOT

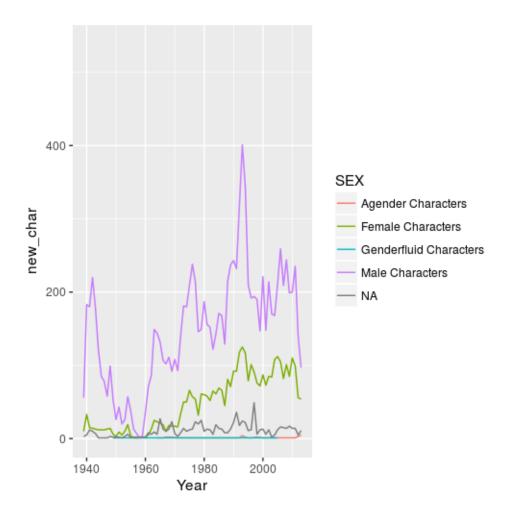


Figure 2.5:

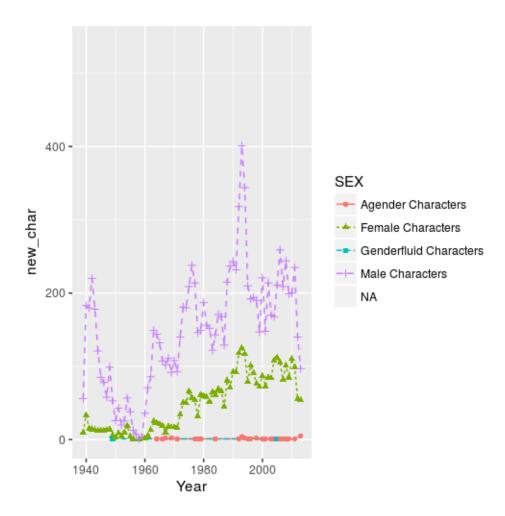


Figure 2.6:

2.3. LINE PLOT 17

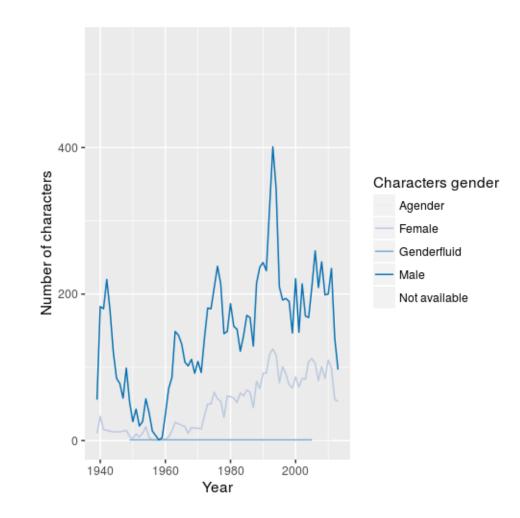


Figure 2.7:

### 2.4 Barplot

Let us consider the marvel\_wikia\_data dataset.

### 2.4.1 Exercise 1

- a. Build a stacked barplot for representing the number of new comic characters distinguishing them by ALIGN and map fill to SEX. Set bars width as 0.7.
- b. advanced: Rotate the x axis by  $30^{\circ}$  so that the axis text nomore overlaps.

```
ggplot(data=marvel_wikia_data, mapping=aes(x=ALIGN, fill=SEX)) +
    geom_bar(width=0.7) +
    theme(axis.text.x = element_text(angle=30, hjust=1, vjust=1))
```

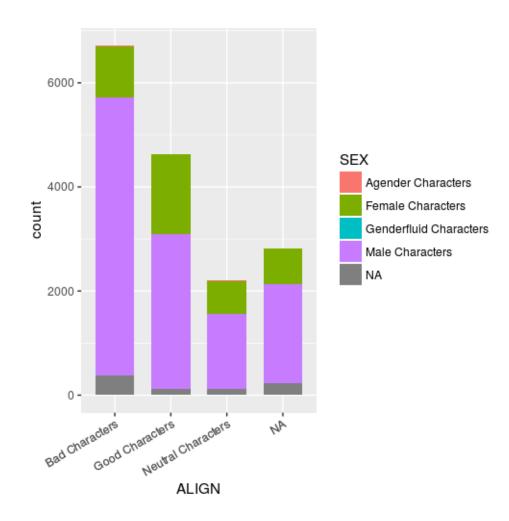


Figure 2.8:

2.4. BARPLOT 19

c. Consider only comic characters with blond hair and Black Hair (filter(HAIR == "Black Hair" | HAIR == "Blond Hair")). Build a stacked barplot for representing the number of new comic characters distinguishing them by ALIGN and map fill to HAIR.

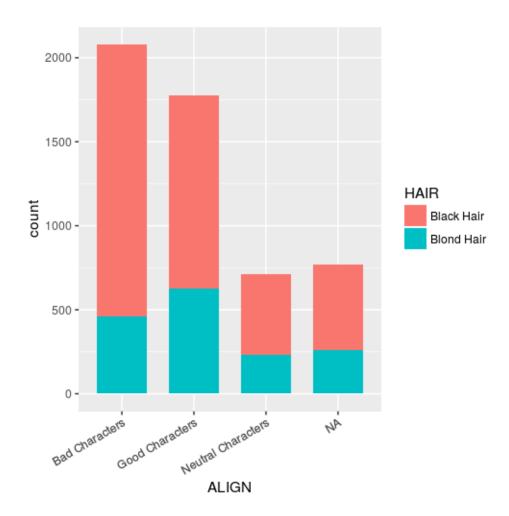


Figure 2.9:

- d. Take the barplot in (b.) and represent the distribution on Blond Hair between the character type (Good, Bad, neutral).
- e. *advanced*: Manually set colour grey20 for black hair and gold3 for blond hair and change the axis name from ALIGN to Character Type.

```
pl + scale_fill_manual(values=c("grey20", "gold3")) +
    labs(x = "Character type")
```

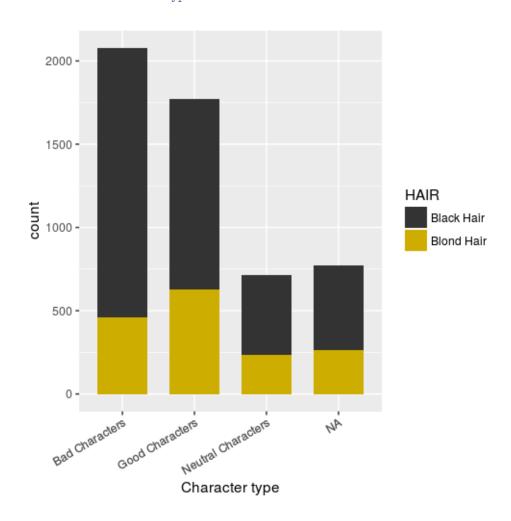


Figure 2.10:

### 2.4.2 Exercise 2

a. Consider only female and male comic characters (filter(SEX == "Male Characters" | SEX == "Female Characters")). Build a barplot with dodged barsfor representing the number comic characters distinguishing them by ALIGN and flip coordinates. Set bars width as 0.5.

2.4. BARPLOT 21

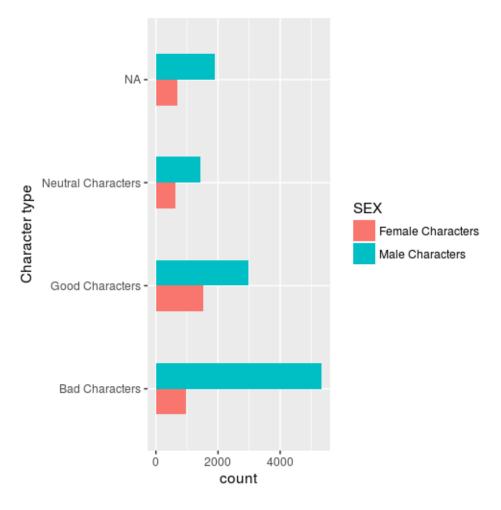


Figure 2.11:

- b. advanced: Consider only comic characters with blue, black and brown eyes. Set facet\_grid(~ EYE)
- c. *advanced*: Customise legend, axis names and colours so that your plot is as clear as possible (for instance, you may choose colour blue for males and pink for females).



Figure 2.12:

2.5. HISTOGRAM 23

### 2.5 Histogram

Let us consider iris dataset.

### **2.5.1** Exercise 1

```
a. Represent the distribution of Sepal_Length variable with an histogram.
```

- b. Set bins fill colour as "hotpink" and bins line colour as "deeppink".
- c. Set the number of bins as 15.

```
pl <- ggplot(data=iris, aes(x=Sepal.Length)) +
    geom_histogram(fill="hotpink", colour="deeppink", bins=15)
pl</pre>
```

d. advanced: Map the grouping variable Species to fill and choose a pink colour palette (PuRd)

```
ggplot(data=iris, aes(x=Sepal.Length, fill = Species)) +
geom_histogram(bins = 14, colour="deeppink") +
scale_fill_brewer(palette="PuRd")
```

e. advanced: Using facet\_grid() produce a different panel for each Species

```
pl +
  facet_grid(Species ~ .)
```

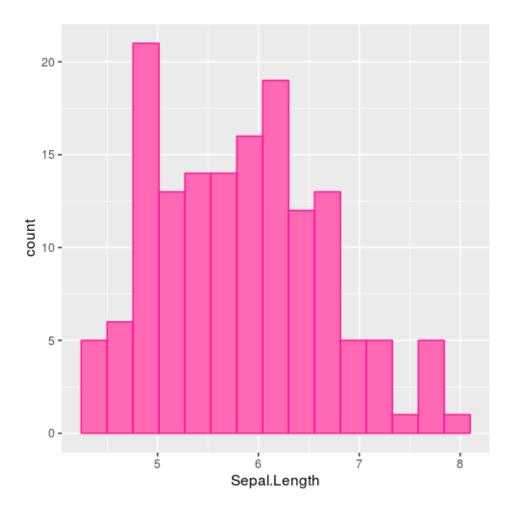


Figure 2.13:

2.5. HISTOGRAM 25

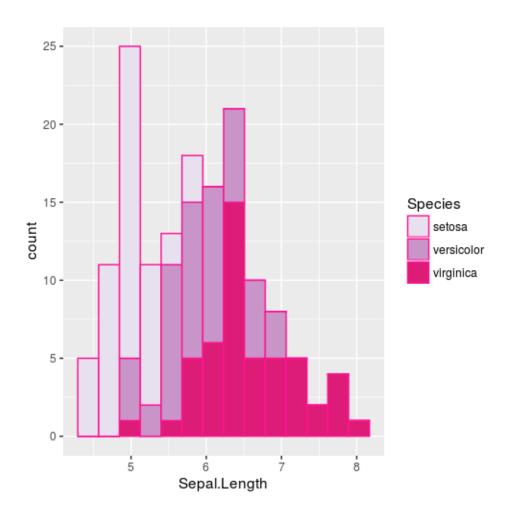


Figure 2.14:

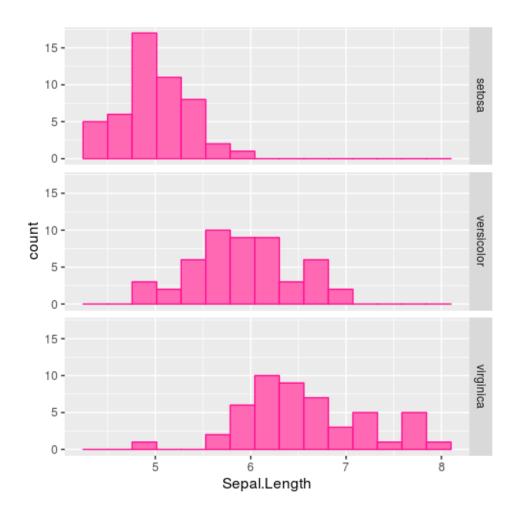


Figure 2.15:

2.6. BOXPLOT 27

## 2.6 Boxplot

### **2.6.1** Exercise 1

a. Build a boxplot to represent the number of times that each comic character created in 2012 have appeared. Highlight outliers in red and set outlier.shape=10 and outlier.size=2. Choose fill = "aquamarine2" and color = "aquamarine4"

```
ggplot(data=marvel_wikia_data %>% filter(Year == 2012), aes(x = 0, y = APPEARANCES)) +
xlab("") + geom_boxplot(colour = "aquamarine4", fill = "aquamarine2",
outlier.colour="red", outlier.shape=10, outlier.size=2)
```

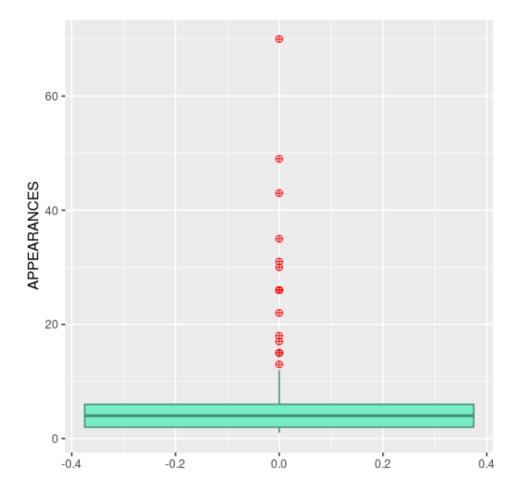


Figure 2.16:

b. Compare the number of times Bad comic characters and Good comic characters created in 2012 have appeared.

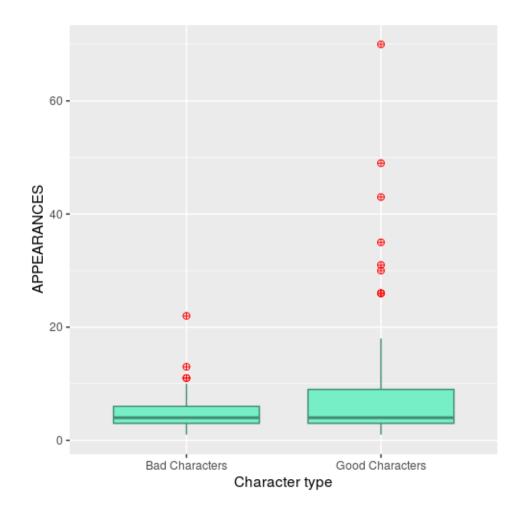


Figure 2.17: