

# Einführung in R: Teil 1

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## Einführung

### 1. Einfache mathematische Operationen

```
4 + 5
```

```
## [1] 9
```

```
sqrt(36)
```

```
## [1] 6
```

```
4 + 2 * 2
```

```
## [1] 8
```

```
log10(100)
```

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

```
log(100, base = 4)
```

```
## [1] 3.321928
```

```
50 + pi
```

```
## [1] 53.14159
```

### 2. Variablen

#### 2.1 Variablen zuweisen

```

x <- 5
a <- 2 + 9
b <- (x + a) * 2
x <- b - 2

alter <- 30
name <- "Max"

mein_vergleich <- 6 > 8
neue_var <- (x < 10 && x > 5) | b == 65

heights <- c(163, 170, NA, 167, NA)

```

### Exercise 0:

Was ist der aktuelle Wert von b?

## 2.2 Datentypen überprüfen

```
class(alter)
```

```
## [1] "numeric"
```

```
class(name)
```

```
## [1] "character"
```

```
class(mein_vergleich)
```

```
## [1] "logical"
```

## 3. Einfache Funktionen

### 3.1 Eingebaute Funktionen

```
heights <- c(163, 170, NA, 167)
mean(heights)
```

```
## [1] NA
```

```
which(is.na(heights))
```

```
## [1] 3
```

```
?mean
```

```
mean(heights, na.rm = TRUE)
```

```
## [1] 166.6667
```

### Exercise 1.1

- a) Was ist die Summe der ersten 100 positiven ganzen Zahlen? Die Formel für die Summe der ganzen Zahlen von 1 bis  $n$  lautet  $\frac{n(n+1)}{2}$ .

```
n <- 100
```

```
n*(n+1)/2
```

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

- b) Was ist die Summe der ersten 1000 positiven ganzen Zahlen?

### Exercise 1.2

Probiere `sum(seq(1,n))` aus. Was macht `seq()`? Was macht `sum()`?

## 3.2 Eigene Funktionen erstellen

```
# Eine einfache Funktion, die zwei Zahlen addiert
addiere <- function(a, b) {
  return(a + b)
}

addiere(5, 7)
```

```
## [1] 12
```

```
numbers_to_n <- function(n) {
  summe <- n*(n+1)/2
}
numbers_to_n(100)
```

## 4. Hilfe in R

```
?seq
vignette(package = "dplyr")
```

## Packages

```
install.packages(c("ggplot2", "dslabs", "dplyr"))
```

## 5. Einfache Navigation in Dataframes

Erkunde den Datensatz “murders”

```
library(dslabs)
data(murders)

print(murders)
```

```
##           state abb      region population total
## 1      Alabama  AL      South    4779736    135
## 2       Alaska  AK      West     710231     19
## 3      Arizona  AZ      West    6392017    232
## 4      Arkansas AR      South    2915918     93
## 5    California CA      West   37253956   1257
## 6      Colorado CO      West    5029196     65
## 7    Connecticut CT    Northeast    3574097     97
## 8      Delaware DE      South     897934     38
## 9 District of Columbia DC      South     601723     99
## 10     Florida  FL      South   19687653   669
## 11     Georgia  GA      South   9920000    376
## 12     Hawaii  HI      West    1360301      7
## 13     Idaho   ID      West    1567582     12
## 14    Illinois IL North Central  12830632   364
## 15     Indiana IN North Central   6483802   142
## 16      Iowa   IA North Central   3046355     21
## 17     Kansas  KS North Central   2853118     63
## 18     Kentucky KY      South   4339367    116
## 19    Louisiana LA      South   4533372    351
## 20      Maine  ME    Northeast   1328361     11
## 21     Maryland MD      South   5773552    293
## 22 Massachusetts MA    Northeast   6547629    118
## 23      Michigan MI North Central   9883640   413
## 24     Minnesota MN North Central   5303925     53
## 25    Mississippi MS      South   2967297    120
## 26     Missouri MO North Central   5988927    321
## 27      Montana MT      West     989415     12
## 28     Nebraska NE North Central   1826341     32
## 29      Nevada  NV      West    2700551     84
## 30    New Hampshire NH    Northeast   1316470      5
## 31     New Jersey NJ    Northeast   8791894   246
## 32     New Mexico NM      West    2059179     67
## 33      New York NY    Northeast  19378102   517
## 34    North Carolina NC      South   9535483   286
## 35     North Dakota ND North Central    672591      4
## 36      Ohio    OH North Central  11536504   310
```

```
## 37      Oklahoma OK      South 3751351 111
## 38      Oregon  OR      West  3831074  36
## 39      Pennsylvania PA    Northeast 12702379 457
## 40      Rhode Island RI    Northeast 1052567 16
## 41      South Carolina SC    South 4625364 207
## 42      South Dakota SD North Central 814180 8
## 43      Tennessee TN      South 6346105 219
## 44      Texas TX      South 25145561 805
## 45      Utah UT      West 2763885 22
## 46      Vermont VT    Northeast 625741 2
## 47      Virginia VA      South 8001024 250
## 48      Washington WA      West 6724540 93
## 49      West Virginia WV    South 1852994 27
## 50      Wisconsin WI North Central 5686986 97
## 51      Wyoming WY      West 563626 5
```

```
head(murders)
```

```
##      state abb region population total
## 1  Alabama AL  South  4779736  135
## 2  Alaska  AK  West   710231   19
## 3  Arizona AZ  West  6392017  232
## 4  Arkansas AR  South 2915918   93
## 5  California CA West 37253956 1257
## 6  Colorado CO  West  5029196   65
```

```
str(murders)
```

```
## 'data.frame': 51 obs. of 5 variables:
## $ state : chr "Alabama" "Alaska" "Arizona" "Arkansas" ...
## $ abb : chr "AL" "AK" "AZ" "AR" ...
## $ region : Factor w/ 4 levels "Northeast","South",...: 2 4 4 2 4 4 1 2 2 2 ...
## $ population: num 4779736 710231 6392017 2915918 37253956 ...
## $ total : num 135 19 232 93 1257 ...
```

```
names(murders)
```

```
## [1] "state" "abb" "region" "population" "total"
```

```
nrow(murders)
```

```
## [1] 51
```

```
table(murders$region)
```

```
##
## Northeast      South North Central      West
##           9           17           12           13
```

## 5.2 Spalten und Zeilen auswählen

```
# Eine Spalte auswählen
murders$state
```

```
## [1] "Alabama"      "Alaska"        "Arizona"
## [4] "Arkansas"     "California"    "Colorado"
## [7] "Connecticut"  "Delaware"      "District of Columbia"
## [10] "Florida"      "Georgia"       "Hawaii"
## [13] "Idaho"        "Illinois"      "Indiana"
## [16] "Iowa"         "Kansas"        "Kentucky"
## [19] "Louisiana"    "Maine"         "Maryland"
## [22] "Massachusetts" "Michigan"      "Minnesota"
## [25] "Mississippi"  "Missouri"      "Montana"
## [28] "Nebraska"     "Nevada"        "New Hampshire"
## [31] "New Jersey"   "New Mexico"    "New York"
## [34] "North Carolina" "North Dakota"  "Ohio"
## [37] "Oklahoma"     "Oregon"        "Pennsylvania"
## [40] "Rhode Island" "South Carolina" "South Dakota"
## [43] "Tennessee"    "Texas"         "Utah"
## [46] "Vermont"      "Virginia"      "Washington"
## [49] "West Virginia" "Wisconsin"     "Wyoming"
```

```
# Eine Zeile auswählen
murders[2, ]
```

```
## state abb region population total
## 2 Alaska AK West 710231 19
```

```
# Eine spezifische Zelle auswählen
murders[3, "population"]
```

```
## [1] 6392017
```

## 5.3 Beispiele der einfachen Manipulationen

```
murders_sued <- gsub("South", "Sued", murders$region)
murders$region <- murders_sued

murders$murder_rate <- murders$total / murders$population
murders$rate_percent <- murders$total * 100 / murders$population

lowest <- min(murders$murder_rate)
highest <- max(murders$murder_rate)

murders_west <- murders[which(murders$region == "West"),]

murders_west[which(murders_west$murder_rate == min(murders_west$murder_rate) |
murders_west$murder_rate == max(murders_west$murder_rate)), "state"]

## [1] "Arizona" "Hawaii"
```

```
murders[which(murders$murder_rate == max(murders$murder_rate)), "state"]
```

```
## [1] "District of Columbia"
```

```
murders_20_100 <- murders[which(murders$total > 20 & murders$total < 100),]
```

```
murders_1000000 <- murders[which(murders$population > 1000000),]
```

```
murders_kleiner1000000 <- murders[which(murders$population <= 1000000),]
```

```
t.test(murders_1000000$murder_rate, murders_kleiner1000000$murder_rate)
```

```
##
```

```
## Welch Two Sample t-test
```

```
##
```

```
## data: murders_1000000$murder_rate and murders_kleiner1000000$murder_rate
```

```
## t = -0.39345, df = 7.1948, p-value = 0.7054
```

```
## alternative hypothesis: true difference in means is not equal to 0
```

```
## 95 percent confidence interval:
```

```
## -5.299978e-05 3.780729e-05
```

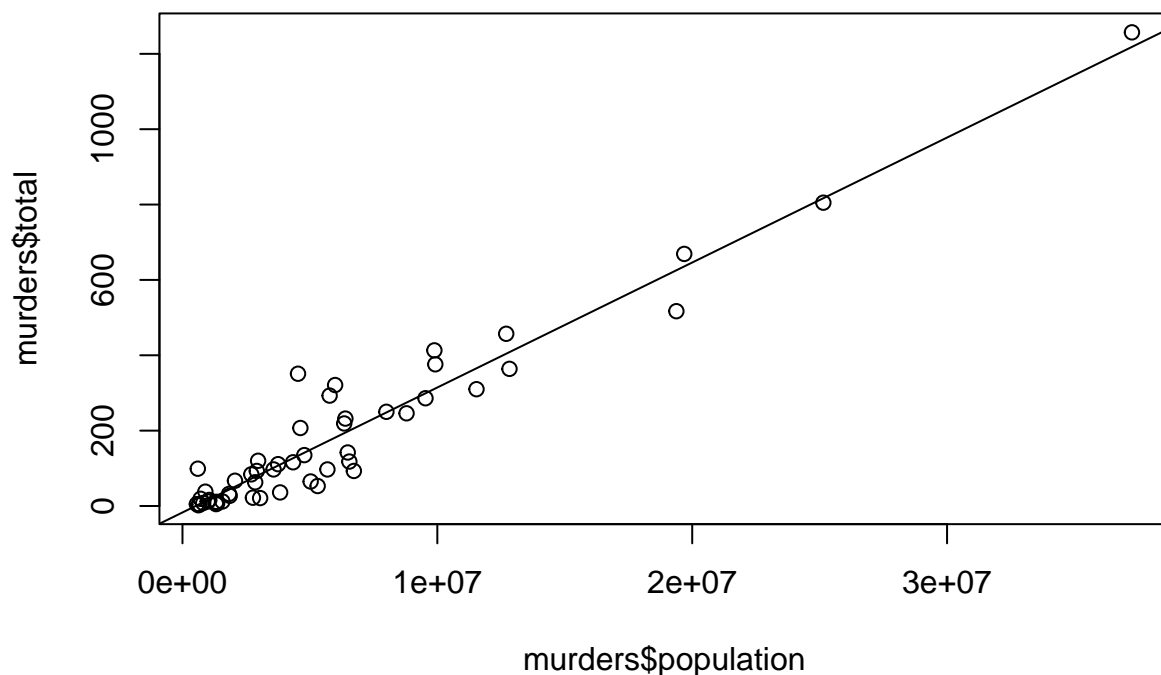
```
## sample estimates:
```

```
## mean of x mean of y
```

```
## 2.659969e-05 3.419593e-05
```

```
plot(murders$population, murders$total)
```

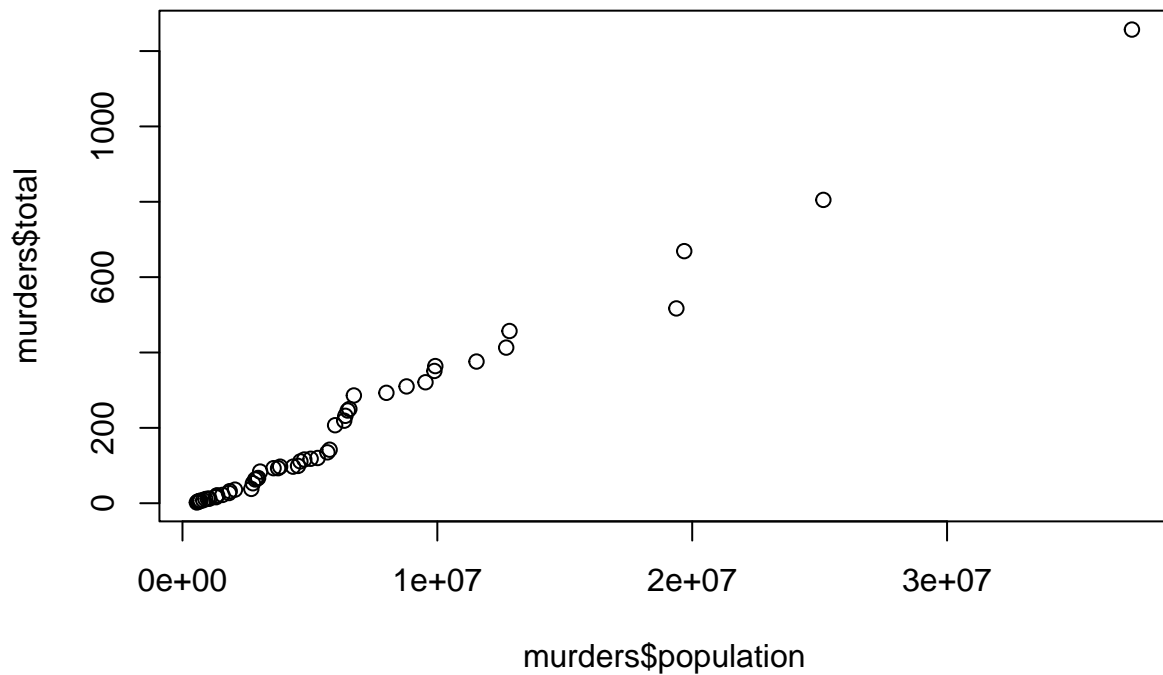
```
abline(lm(total ~ population, murders))
```



```
cor(murders$population, murders$total)
```

```
## [1] 0.9635956
```

```
qqplot(murders$population, murders$total)
```



# 6. Wrangling mit dplyr

```
library(dplyr)
```

```
data(mtcars)
```

```
data <- mtcars
```

```
data$auto_marke <- rownames(data)
```

```
rownames(data) <- NULL
```

```
# write.table(data, "output_data/mtcars.txt", sep = "\t")
```

```
selected_data <- select(data, mpg, cyl, hp, auto_marke)
```

```
filtered_data <- filter(data, mpg > 20)
```

```
arranged_data <- arrange(data, desc(mpg))
```



```

mutated_data <- mutate(data, hp_to_weight = hp / wt)

summary_data <- summarise(data, avg_mpg = mean(mpg),
                           avg_hp = median(hp))

# mean over all columns
summary_data_all <- data %>% summarise(across(where(is.numeric), mean, na.rm = TRUE))

summary_data_piped <- data %>%
  filter(mpg > 20) %>%
  mutate(hp_to_weight = hp / wt) %>%
  summarise(avg_hp = mean(hp))

grouped_summary <- data %>%
  group_by(cyl) %>%
  summarise(
    avg_mpg = mean(mpg),
    avg_hp = mean(hp)
  )

#####

df1 <- data.frame(id = 1:5, value1 = letters[1:5])
df2 <- data.frame(id = 3:7, value2 = LETTERS[3:7])

inner <- inner_join(df1, df2, by = "id")

left <- left_join(df1, df2, by = "id")

right <- right_join(df1, df2, by = "id")

full <- full_join(df1, df2, by = "id")

```

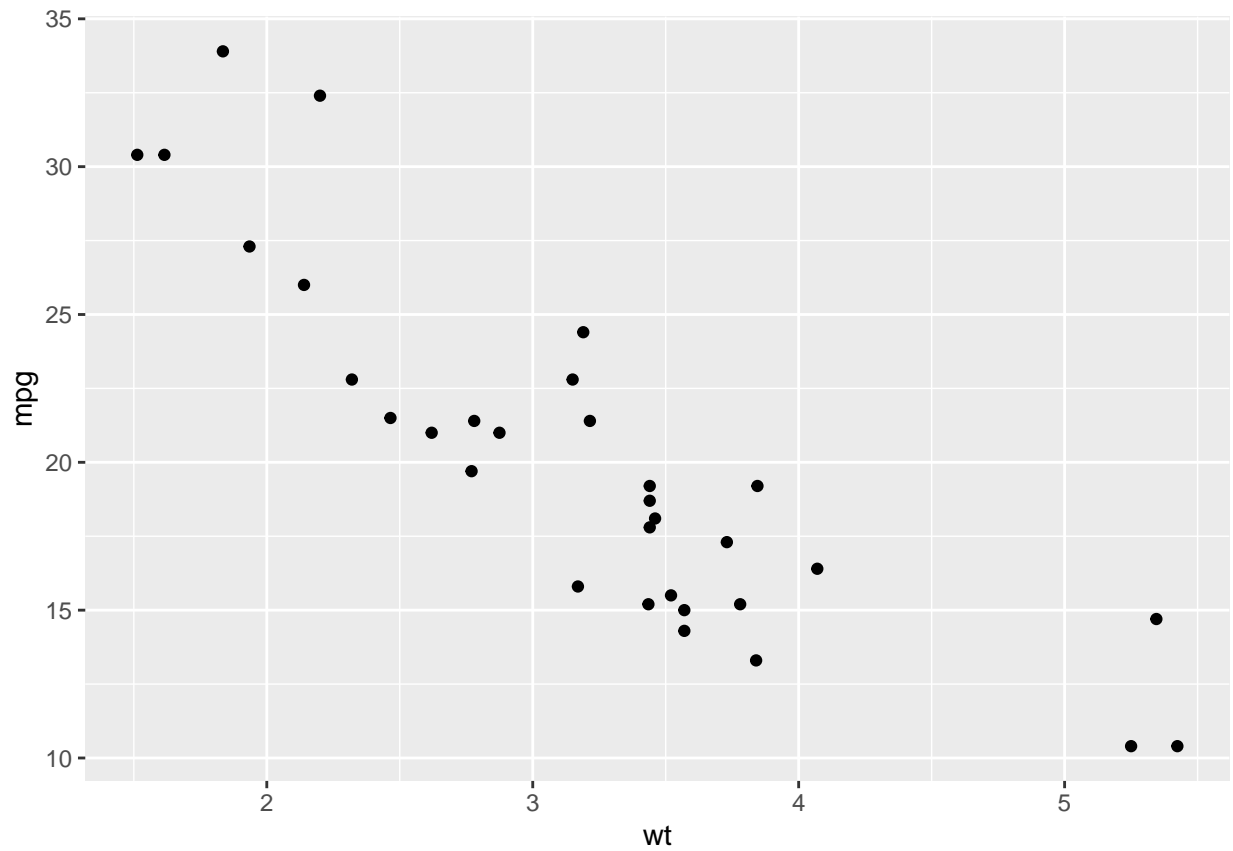
## 7. Daten-Visualisierung

```

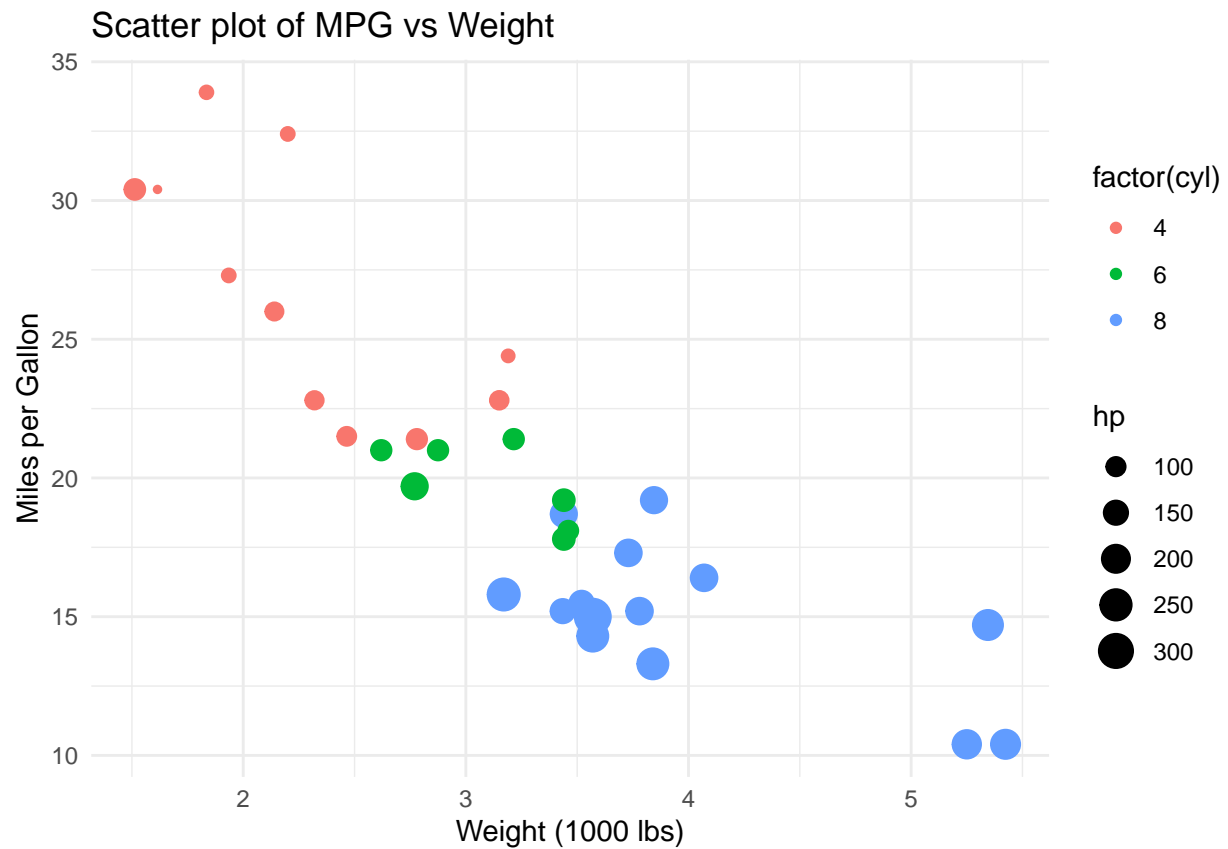
library(ggplot2)

ggplot(mtcars, aes(x = wt, y = mpg)) +
  geom_point()

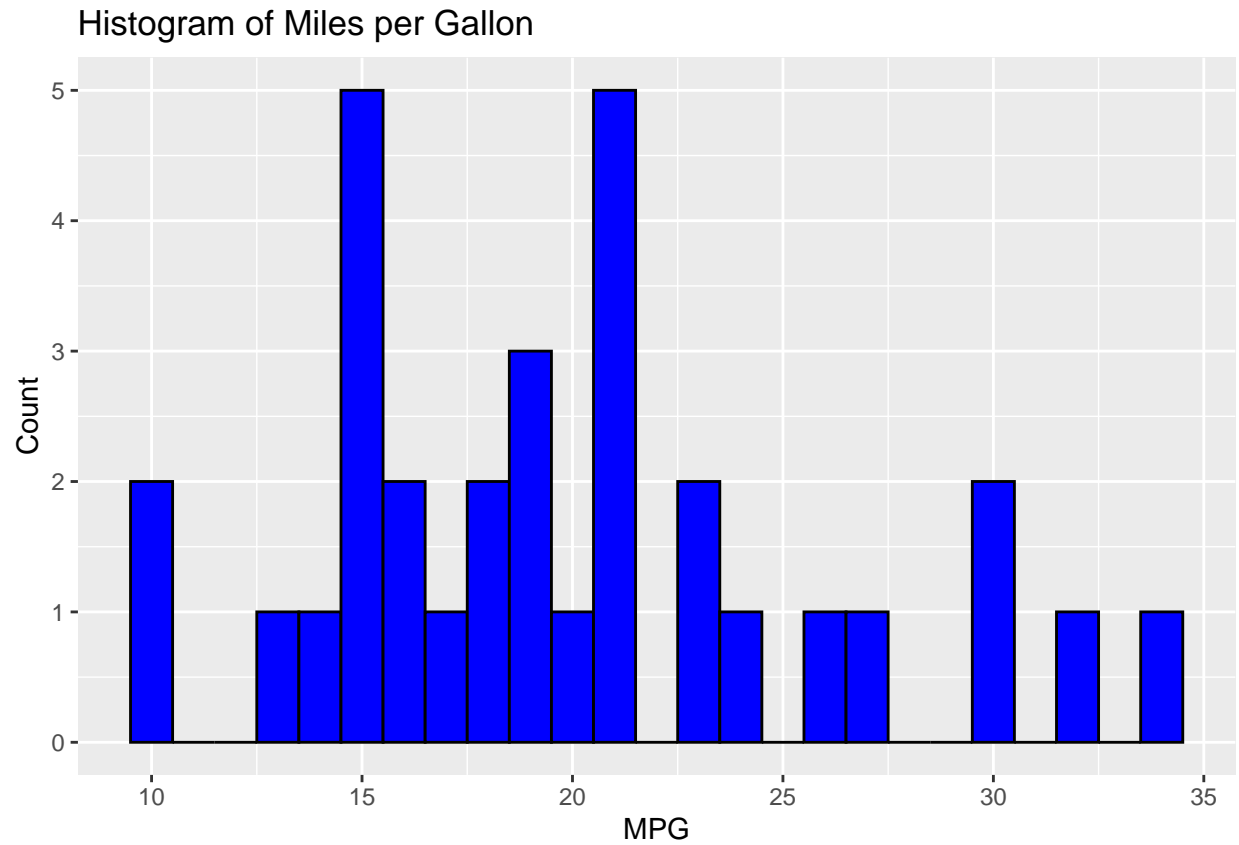
```



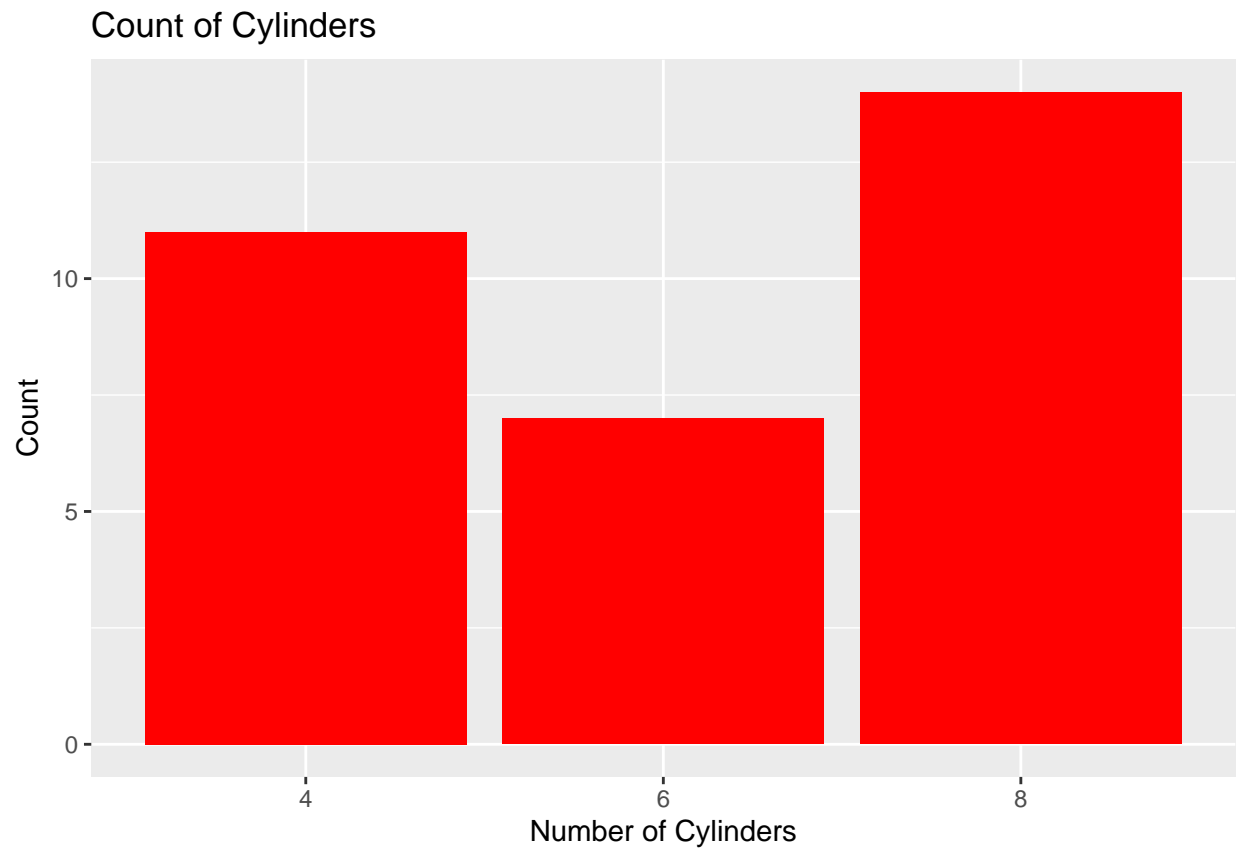
```
ggplot(mtcars, aes(x = wt, y = mpg, color = factor(cyl), size = hp)) +  
  geom_point() +  
  labs(title = "Scatter plot of MPG vs Weight", x = "Weight (1000 lbs)", y = "Miles per Gallon") +  
  theme_minimal()
```



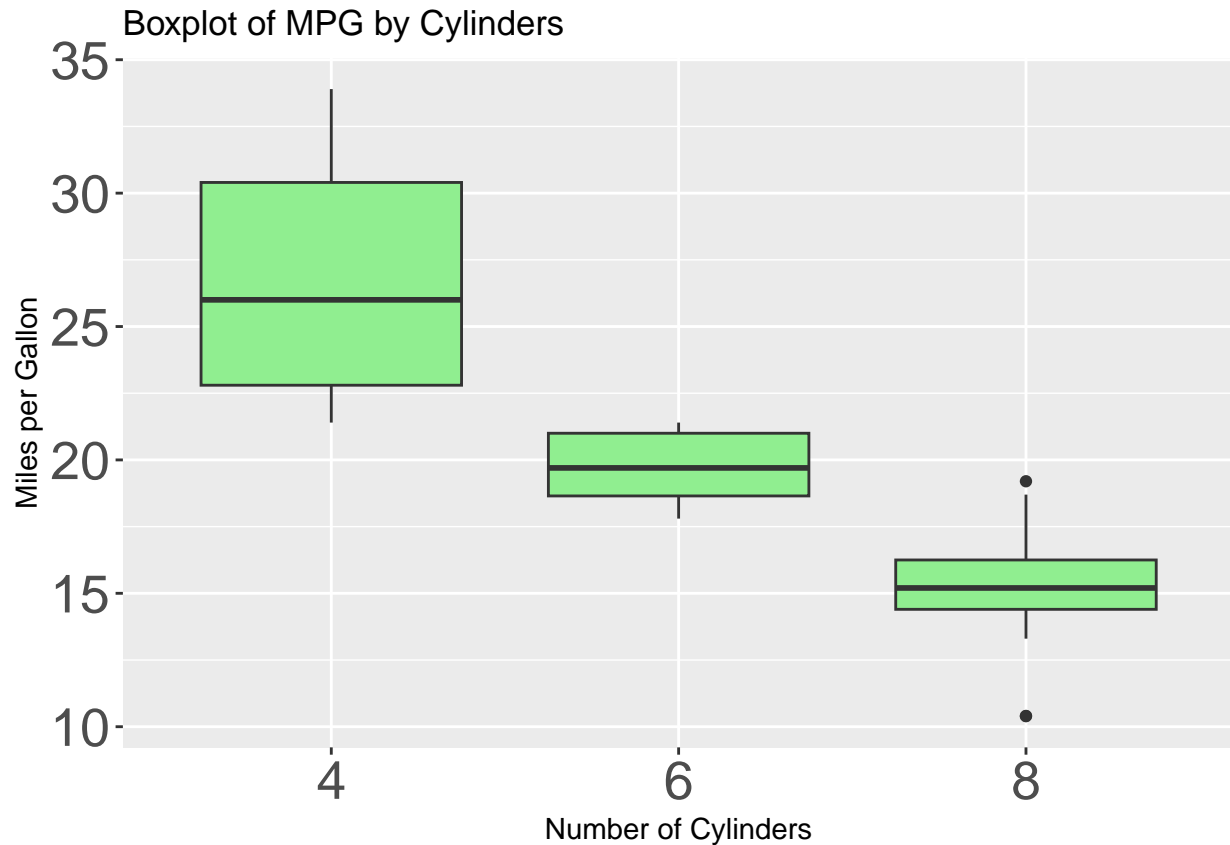
```
ggplot(mtcars, aes(x = mpg)) +  
  geom_histogram(binwidth = 1, fill = "blue", color = "black") +  
  labs(title = "Histogram of Miles per Gallon", x = "MPG", y = "Count")
```



```
ggplot(mtcars, aes(x = factor(cyl))) +  
  geom_bar(fill = "#FF0000") +  
  labs(title = "Count of Cylinders", x = "Number of Cylinders", y = "Count")
```



```
mtcars %>% ggplot(aes(x = factor(cyl), y = mpg)) +  
  geom_boxplot(fill = "lightgreen") +  
  labs(title = "Boxplot of MPG by Cylinders", x = "Number of Cylinders", y = "Miles per Gallon") +  
  theme(axis.text = element_text(size = 20))
```



Farbenpaletten in R [http://www.cookbook-r.com/Graphs/Colors\\_\(ggplot2\)/#a-colorblind-friendly-palette](http://www.cookbook-r.com/Graphs/Colors_(ggplot2)/#a-colorblind-friendly-palette)

## 8. Übungen

### Übung 1

1. Tabelle "survey\_example.txt" einlesen.
2. Mittelwert von height\_cm für männliche und für weibliche StudienteilnehmerInnen ausrechnen.
3. Fehlerquelle identifizieren.
4. Verwendet gsub(), um das Problem zu lösen. Tipp: "[0-9]+" steht für beliebige Anzahl an Zahlen, "\1" steht stellvertretend für eine Zahlengruppe.
5. Zeichnet ein Boxplot mit den Körpergrößen der Männer und der Frauen.

Lösung:

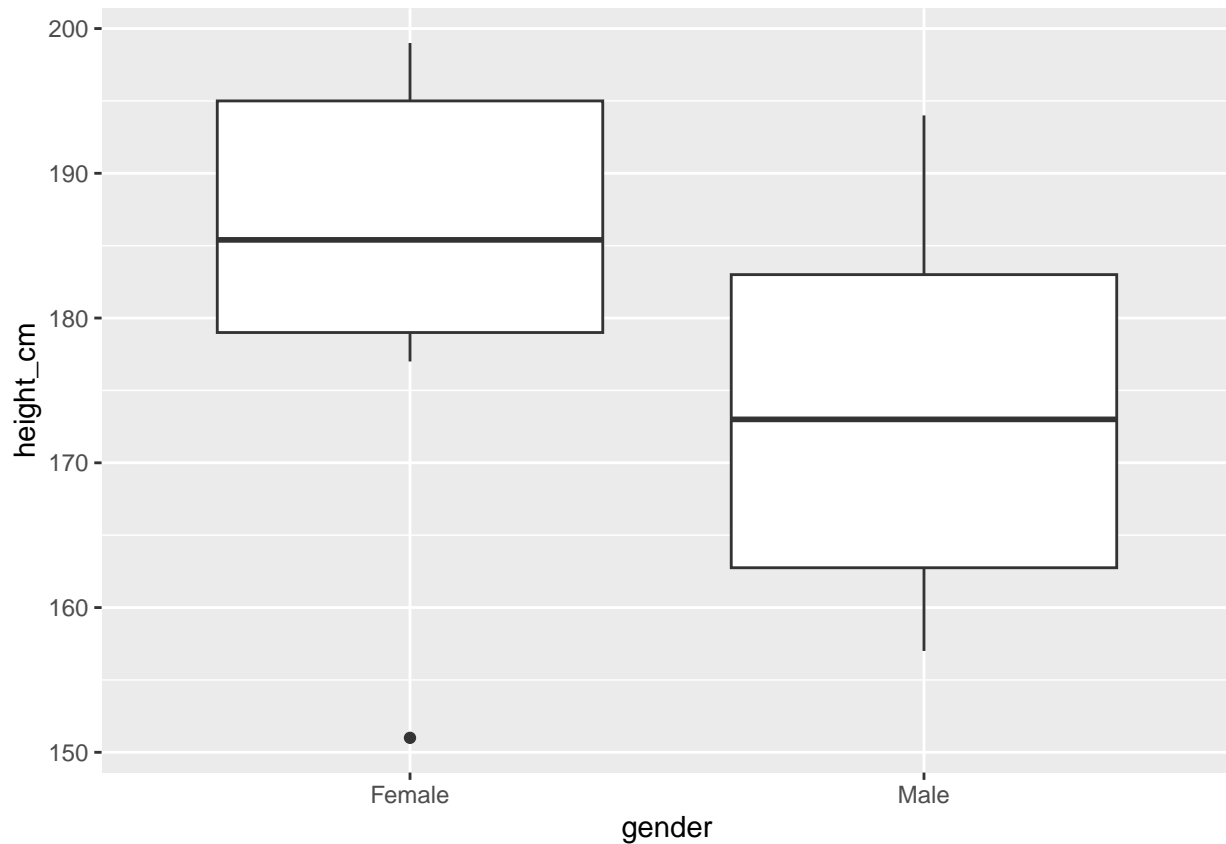
```
heights <- read.table("survey_example.txt")
new_heights <- gsub("[0-9]+cm", "\\1", heights$height_cm)
new_heights <- gsub("[0-9]+", "[0-9]+", "\\1.\\2", new_heights)

heights %>% mutate(height_cm = as.numeric(new_heights)) %>%
  group_by(gender) %>% summarise(avg = mean(height_cm))
```

```
## # A tibble: 2 x 2
##   gender  avg
```

```
##   <chr>  <dbl>
## 1 Female  184.
## 2 Male    174.
```

```
heights %>% mutate(height_cm = as.numeric(new_heights)) %>%
  ggplot(aes(x = gender, y = height_cm)) +
  geom_boxplot()
```



## Übung 2

1. Erkundet den eingebauten Datensatz `data(sleep)`.
2. Ändert die Gruppenbezeichnungen von 1/2 zu "received\_drug\_a"/"received\_drug\_b". Tipp: ihr könnt `recode()` innerhalb von `mutate()` anwenden.
3. Berechnet den Mittelwert von `extra_sleep` für jede Gruppe.
4. Zeigt eure Ergebnisse in einem bar plot. Tipp: verwendet "stat = "identity" Argument.

Lösung:

```
data(sleep)
?slee

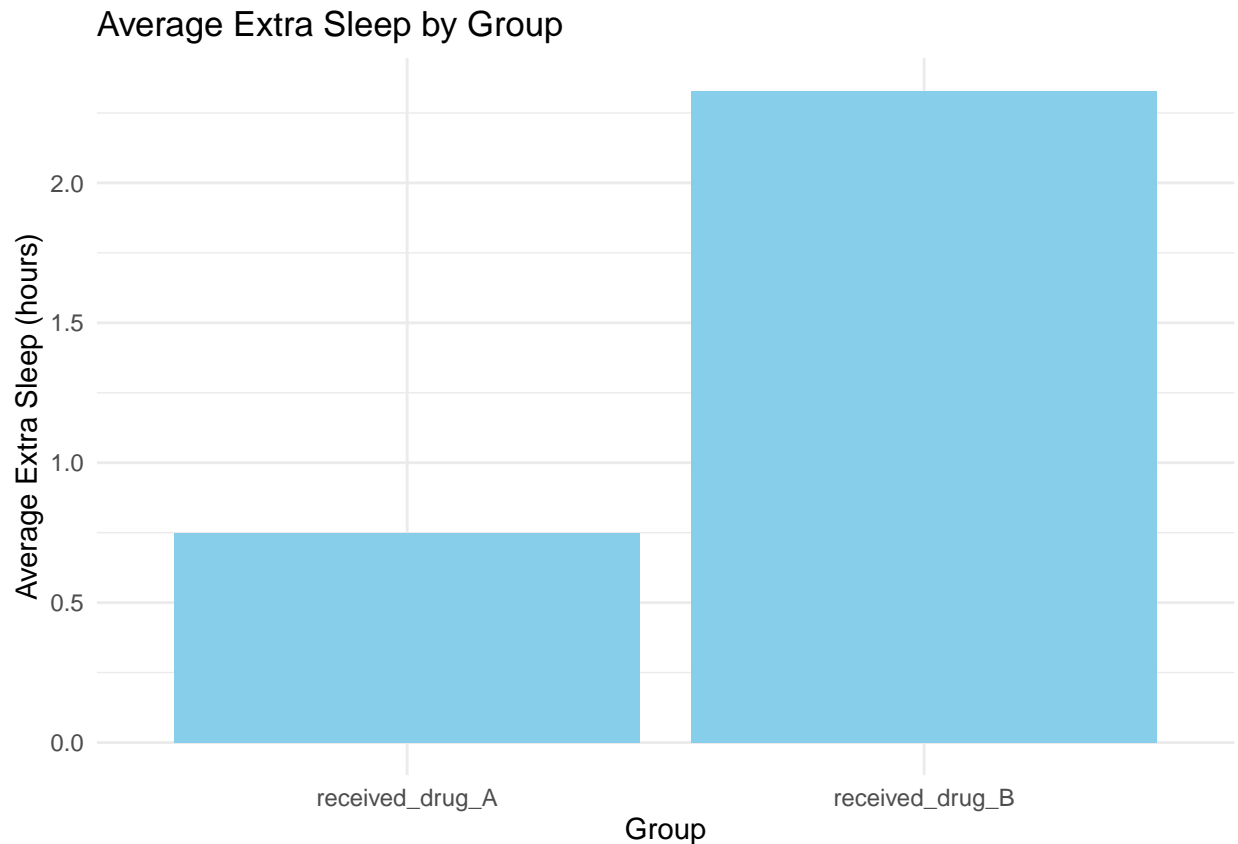
average_sleep <- sleep %>%
  mutate(group = recode(group, `1` = "received_drug_A", `2` = "received_drug_B")) %>%
  group_by(group) %>%
```

```

summarise(mean_sleep = mean(extra))

ggplot(average_sleep, aes(x = factor(group), y = mean_sleep)) +
  geom_bar(stat = "identity", fill = "skyblue") +
  labs(x = "Group", y = "Average Extra Sleep (hours)", title = "Average Extra Sleep by Group") +
  theme_minimal()

```



### Übung 3

0. library(dslabs).
1. Installiert package "ggrepel".
2. Erkundet den eingebauten Datensatz data(divorce\_margarine).
3. Gibt es eine Korrelation zwischen der Anzahl der Scheidungen und dem Margarinekonsum?
4. Versucht die Problematik in einem Bild/Scatter plot/... darzustellen. Tipp: Schaut euch "geom\_text\_repel" function an.

Lösung:

```
install.packages("ggrepel")
```

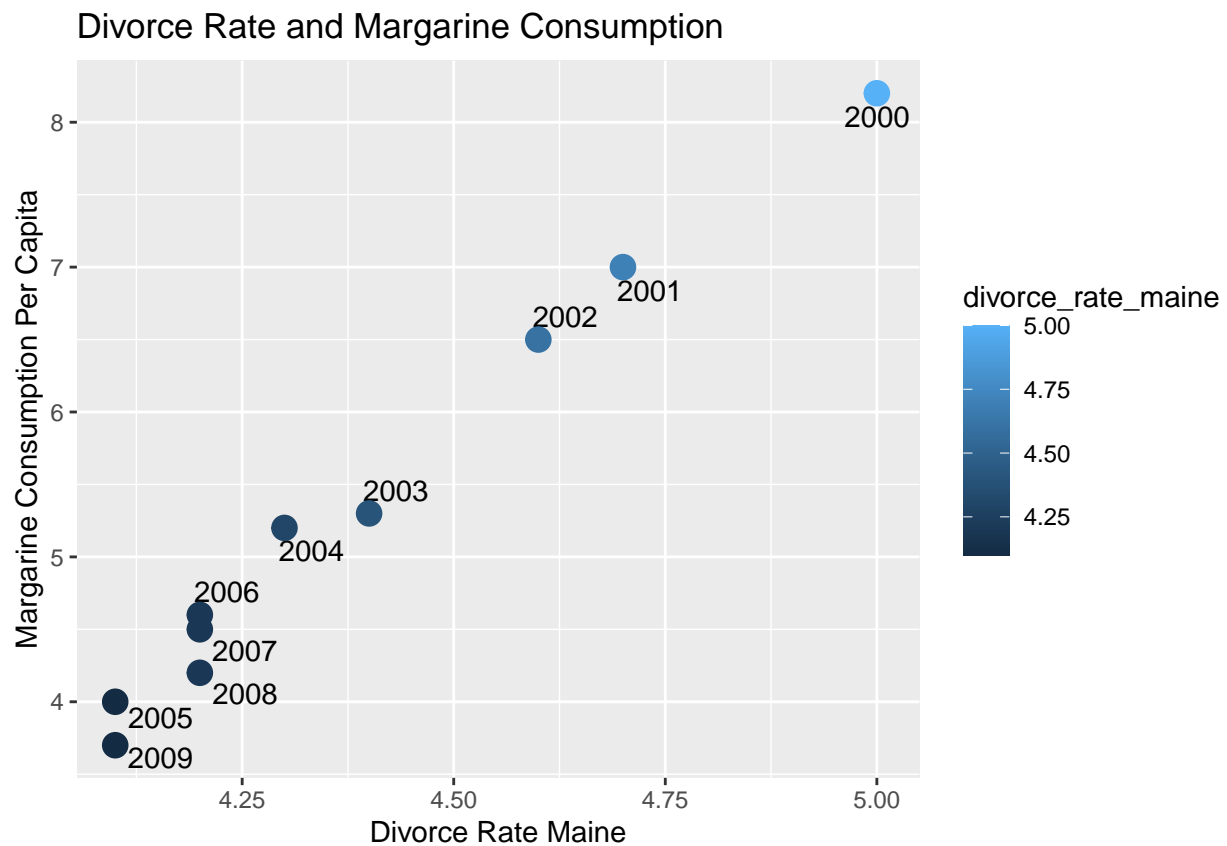
```
install.packages("ggrepel")
```

```
## Installiere Paket nach '/home/ekaterina/R/x86_64-pc-linux-gnu-library/4.1'
## (da 'lib' nicht spezifiziert)
```



```
library(dslabs)
library(ggrepel)
data(divorce_margarine)

divorce_margarine %>%
  ggplot(aes(x = divorce_rate_maine, y = margarine_consumption_per_capita, label = year)) +
  geom_text_repel(nudge_x = 0.005) +
  geom_point(aes(color=divorce_rate_maine), size = 4) +
  xlab("Divorce Rate Maine") +
  ylab("Margarine Consumption Per Capita") +
  ggtitle("Divorce Rate and Margarine Consumption")
```



## Übung 4

1. Erstellt einen "toy"-Datensatz:

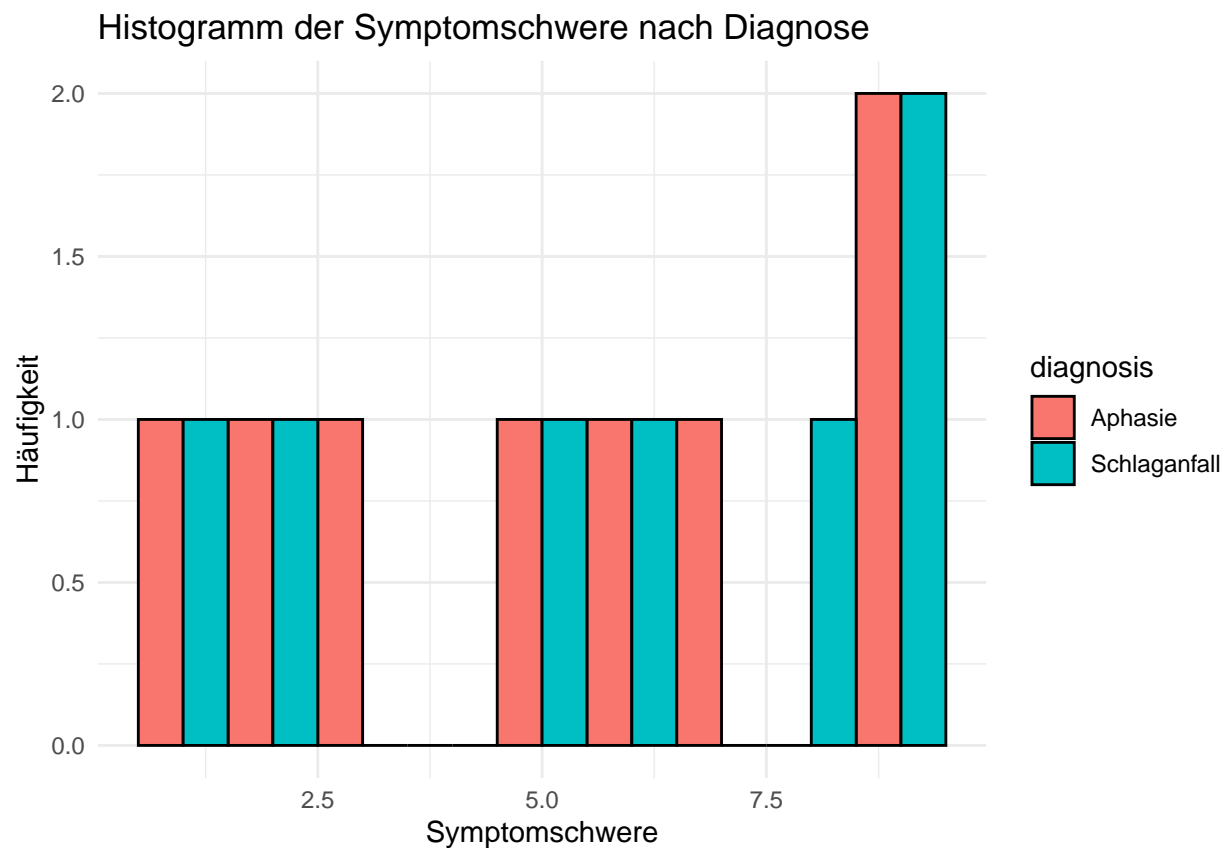
```
neuro_data <- data.frame(
  patient_id = 1:100,
  diagnosis = sample(c("Aphasie", "Dysarthrie", "Schlaganfall"), 100, replace = TRUE),
  symptom_severity = sample(1:10, 100, replace = TRUE),
  therapy_sessions = sample(1:30, 100, replace = TRUE)
)
```

2. Filtriert für Patienten mit Schlaganfall oder Aphasie und mehr als 20 Therapie-Sitzungen.

3. Erstellt ein Histogramm der Symptomschwere.

Lösung:

```
filtered_data <- neuro_data %>%  
  filter(diagnosis %in% c("Aphasie", "Schlaganfall"),  
         therapy_sessions > 20)  
  
ggplot(filtered_data, aes(x = symptom_severity, fill = diagnosis)) +  
  geom_histogram(binwidth = 1, position = "dodge", color = "black") +  
  labs(x = "Symptomschwere", y = "Häufigkeit", title = "Histogramm der Symptomschwere nach Diagnose") +  
  theme_minimal()
```



## Übung 5

1. Führt folgende Befehle aus:

```
install.packages(c("maps", "mapdata"))
```

```
library(ggplot2)  
library(maps)  
library(mapdata)
```

```
# Beispiel Sehenswürdigkeiten in Wien, Salzburg und Graz
```

```
landmarks <- data.frame(
  name = c("Stephansdom, Wien", "Festung Hohensalzburg, Salzburg", "Schlossberg, Graz"),
  lat = c(48.2082, 47.7990, 47.0702),
  lon = c(16.3738, 13.0430, 15.4395)
)

# Landkarte von Österreich
au_map <- map_data("worldHires", region = "Austria")
```

2. Erkundet ggmap-Dokumentation
3. Vervollständigt den Code:

```
ggplot() +
  geom_map(data = XXX, map = XXX,
    aes(x = long, y = lat, map_id = region),
    fill = "lightgrey", color = XXX) +
  geom_point(data = landmarks, aes(x = XXX, y = XXX), XXX = "red", size = 3) +
  geom_text(data = landmarks, aes(x = XXX, y = XXX, label = name), vjust = -1, hjust = 1.5, color = "black") +
  labs(title = "Landmarks in Vienna", x = "Longitude", y = "Latitude") +
  coord_fixed(ratio = 1.2)
```

Lösung:

```
ggplot() +
  geom_map(data = au_map, map = au_map,
    aes(x = long, y = lat, map_id = region),
    fill = "lightgrey", color = "black") +
  geom_point(data = landmarks, aes(x = lon, y = lat), color = "red", size = 3) +
  geom_text(data = landmarks, aes(x = lon, y = lat, label = name), vjust = -1, hjust = 1.5, color = "black") +
  labs(title = "Landmarks in Vienna", x = "Longitude", y = "Latitude") +
  coord_fixed(ratio = 1.2)
```

```
## Warning in geom_map(data = au_map, map = au_map, aes(x = long, y = lat, :
## Ignoring unknown aesthetics: x and y
```

## Landmarks in Vienna

