

`— title: "ExamPrep" output: pdf_document —

От упражнение 1

```
#1.
#create a vector
vect <- c(8,3,8,7,15,9,12,4,9,10,5,1)
#create a 4x3 matrix
m <- matrix(vect, nrow = 4, ncol = 3)
#adding a column
m1 <- cbind(m, c(1,3,5,7))

#indexes of the first column
ordered <- order(m1[,1], decreasing = FALSE)
#ordered matrix
ordered_by_first_column <- m1[ordered,]

#indexes of the first two columns
ordered2 <- order(m1[,1], m1[,2], decreasing = FALSE)
#ordered matrix
ordered_by_two_columns <- m1[ordered2, ]

#2.
#most and least expensive in 2000
most <- which.max(homedata$y2000)
least <- which.min(homedata$y2000)

#prices in 1970
homedata$y1970[most]

## [1] 198900

homedata$y1970[least]

## [1] 10000

#top five most expensive houses in 2000
ordered_five <- homedata$y2000[order(homedata$y2000, decreasing = T)]

top_five <- head(ordered_five ,5)

#средната цена на 5те най-скъпи от 2000, но на техните цени от 1970
mean_top_five <- mean(head(homedata$y1970[order(homedata$y2000, decreasing = T)], 5))

#къщите, чийто цена е намаляла през 2000г.
lowered <- homedata$y2000[which(homedata$y2000 < homedata$y1970)]

percent_increase <- head(order(((homedata$y2000 - homedata$y1970) /
homedata$y1970),
```

```

decreasing = T), 10)

top_ten_increase <- homedata$y2000[percent_increase]

#3.
#number of men
num_men <- nrow(survey[survey$Sex == 'Male', ])

#number of men smokers
num_men_smokers <- nrow(survey[survey$Sex == 'Male' & survey$Smoke !=
'Never', ])

#mean height of all men
mean(survey$Height[survey$Sex == 'Male'], na.rm = T)

## [1] 178.826

#height and sex of the top 6 youngest students
youngest <- head(order(survey$Age), 6)

survey$Sex[youngest]

## [1] Male   Male   Female Female Female Female
## Levels: Female Male

survey$Height[youngest]

## [1]      NA      NA      NA 160.00 172.00 170.18

```

От упражнение 2

```

#1.
#случайно избран човек да се окаже пушач
survey$Smoke %>% table() %>% prop.table()

## .
##      Heavy      Never      Occas      Regul
## 0.04661017 0.80084746 0.08050847 0.07203390

#случайно избран мъж да се окаже редовно пушещ
table(survey$Smoke, survey$Sex) %>% prop.table()

##
##      Female      Male
## Heavy 0.02127660 0.02553191
## Never 0.42127660 0.37872340
## Occas 0.03829787 0.04255319
## Regul 0.02127660 0.05106383

#the same as
smoking_men <- nrow(survey[survey$Sex == 'Male' & survey$Smoke == 'Regul', ])
smoking_men / nrow(survey)

```

```
## [1] 0.05485232
```

#случаен мъж да се окаже редовен пушач. Стойността на всяка клетка се дели на сумата

#от редовете

```
prop.table(table(survey$Sex, survey$Smoke), 1)
```

```
##
```

```
##           Heavy      Never      Occas      Regul
## Female 0.04237288 0.83898305 0.07627119 0.04237288
## Male   0.05128205 0.76068376 0.08547009 0.10256410
```

#случаен редовен пушач да се окаже мъж. Клетката се дели на сумата от колоните

```
prop.table(table(survey$Sex, survey$Smoke), 2)
```

```
##
```

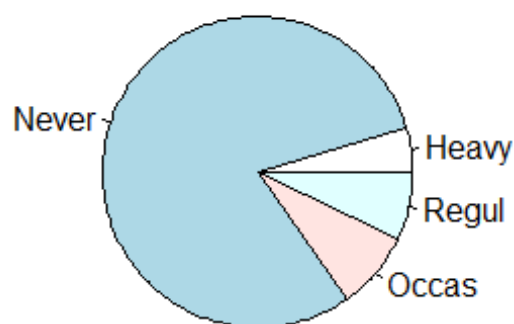
```
##           Heavy      Never      Occas      Regul
## Female 0.4545455 0.5265957 0.4736842 0.2941176
## Male   0.5454545 0.4734043 0.5263158 0.7058824
```

#2.

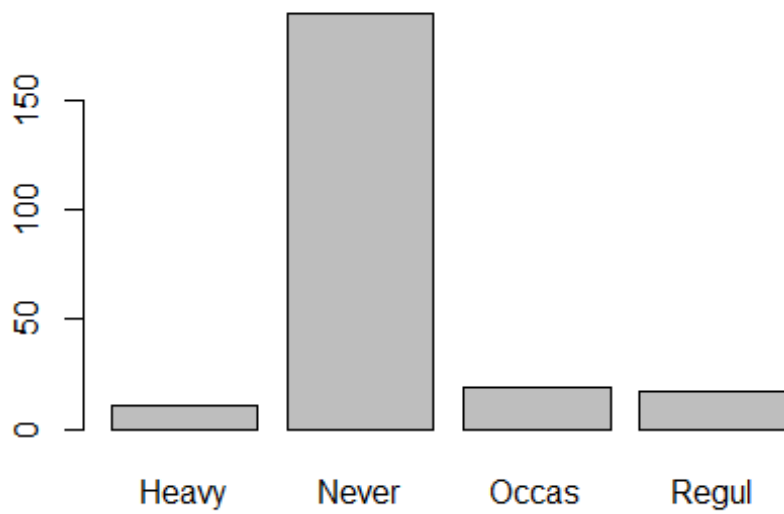
#направете графики за пушачите и за пола

#графики за пушенето

```
pie(table(survey$Smoke))
```

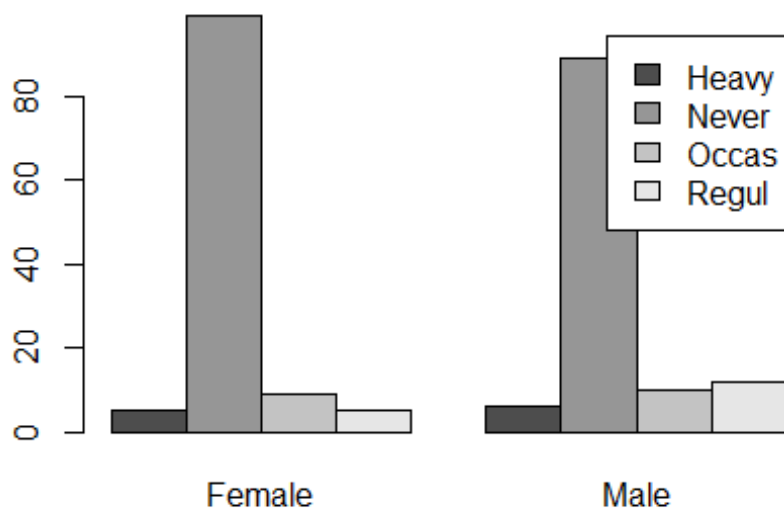


```
barplot(table(survey$Smoke))
```



#графика за пушенето и пола

```
barplot(table(survey$Smoke, survey$Sex), beside = T, legend = T)
```



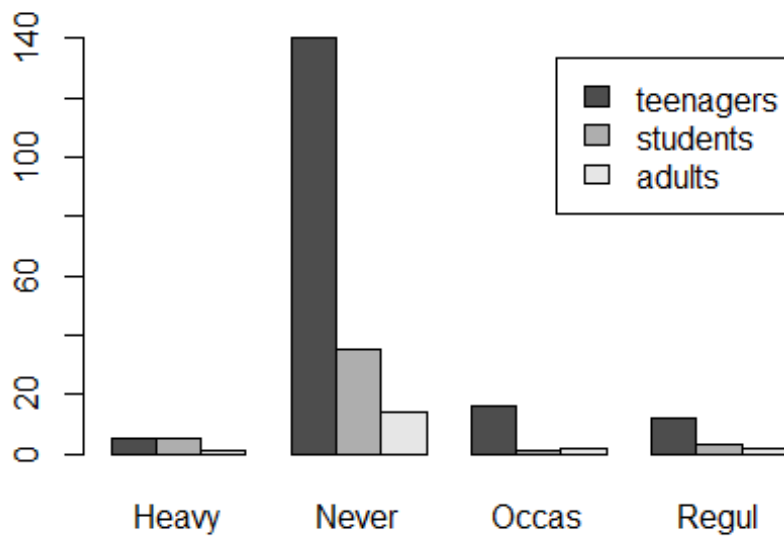
#3.

#за да разделим някаква информация на интервали, които ние искаме ползваме cut

```
groups <- cut(survey$Age, c(0, 20, 25, 100), c('teenagers', 'students',  
'adults'))
```

#правим го на графика

```
table(groups, survey$Smoke) %>%  
  barplot(legend = T, beside = T)
```



#4.

```
s <- sd(survey$Height, na.rm = T)
```

```
med <- median(survey$Height, na.rm = T)
```

```
m <- mean(survey$Height, na.rm = T)
```

```
quantile(survey$Height, na.rm = T)
```

```
## 0% 25% 50% 75% 100%
```

```
## 150 165 171 180 200
```

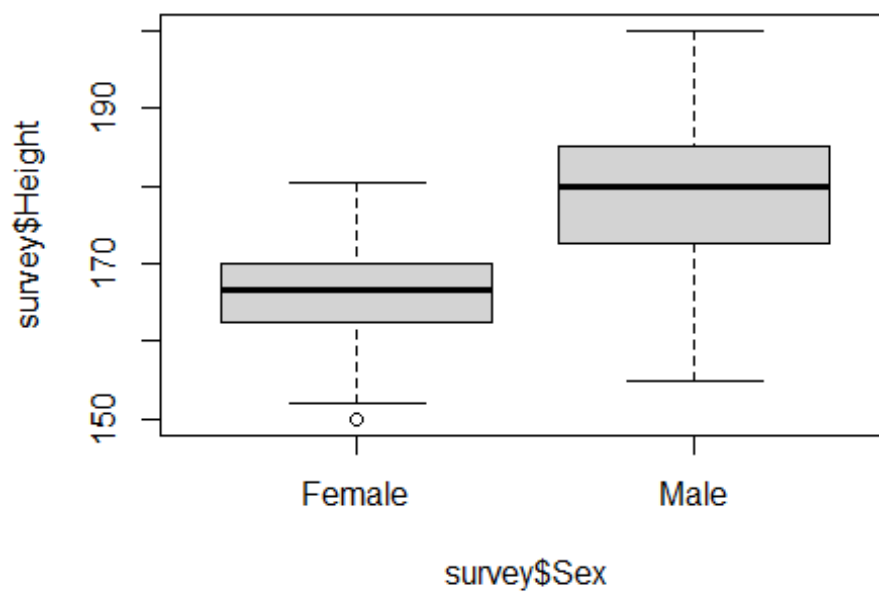
#брой различаващи се от средната височина с не повече от 2 стандартни отклонения

```
cut(survey$Height, c(0, m - s, m + s, 300)) %>%  
  table()
```

```
## .  
## (0,163] (163,182] (182,300]  
##      28      143      38
```

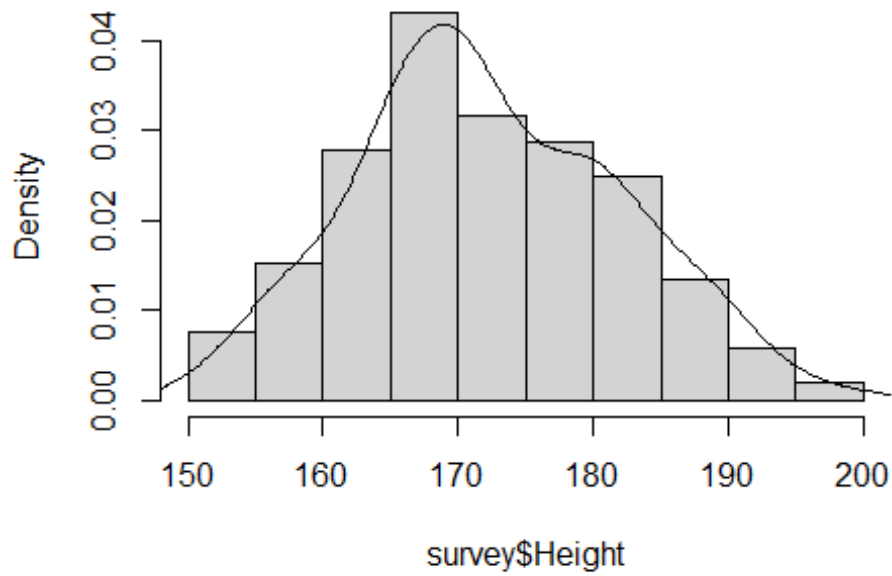
От упражнение 3

```
#1.  
#графика според височината и пола  
boxplot(survey$Height ~ survey$Sex)
```



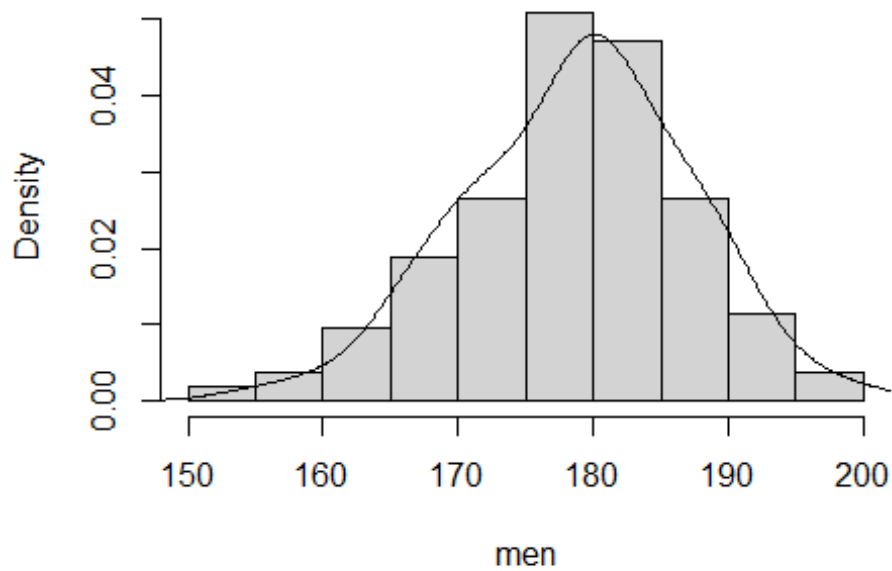
```
#хистограма според височината и имаме плътността  
hist(survey$Height, probability = T)  
lines(density(survey$Height, na.rm = T))
```

Histogram of survey\$Height



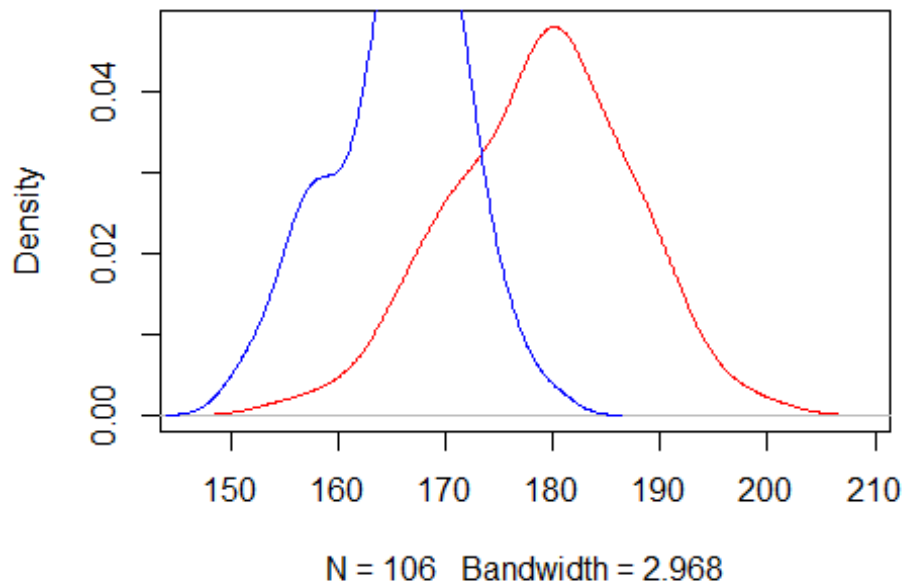
```
#хистограма според височината на мъжете и имаме линия за плътност  
men <- survey$Height[survey$Sex == 'Male']  
women <- survey$Height[survey$Sex == 'Female']  
  
hist(men, probability = T)  
lines(density(men, na.rm = T))
```

Histogram of men

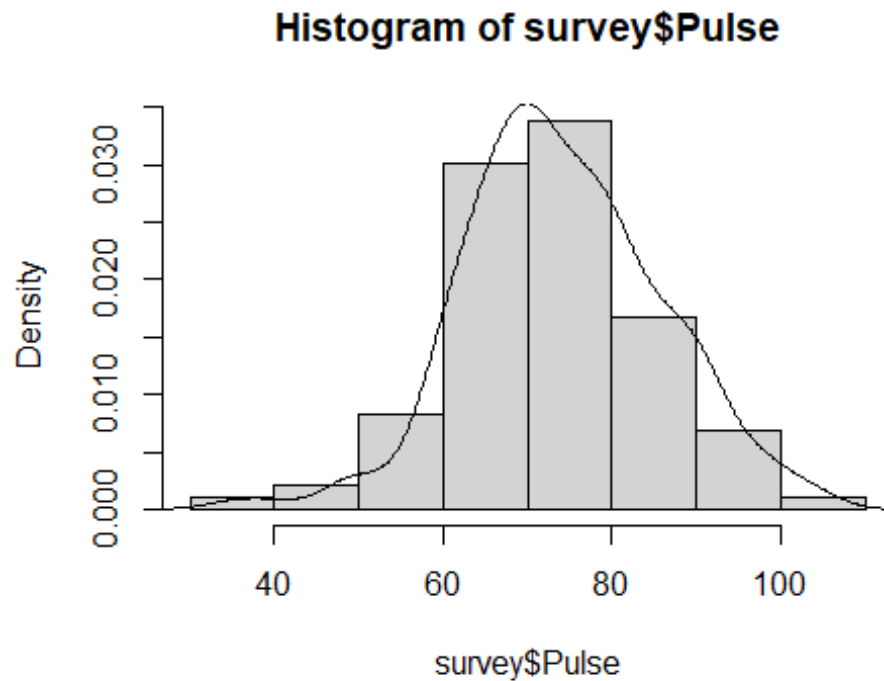


```
#графика за плътностите на височините на двата пола  
plot(density(men, na.rm = T), col='red')  
lines(density(women, na.rm = T), col='blue')
```

density.default(x = men, na.rm = T)

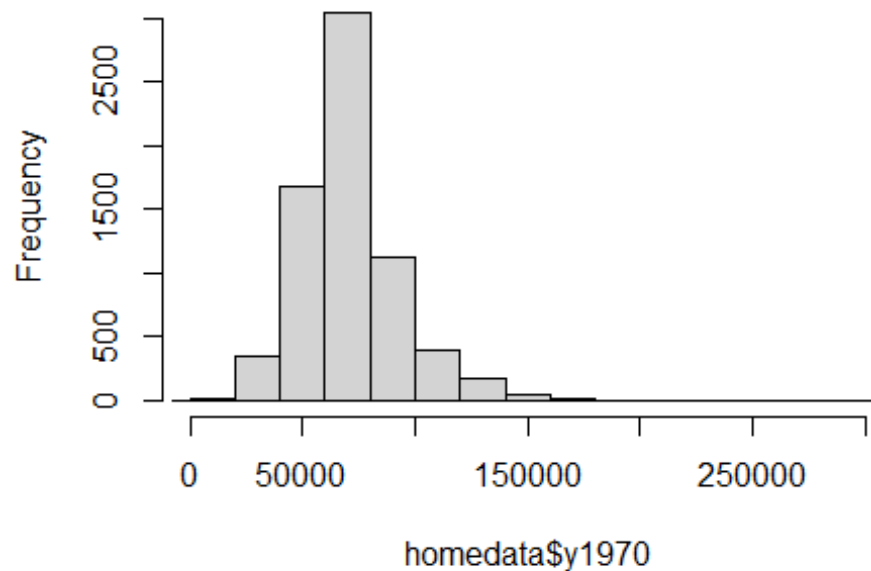



```
#2. Histogram for the pulse of the students including the density  
hist(survey$Pulse, probability = T)  
lines(density(survey$Pulse, na.rm = T))
```



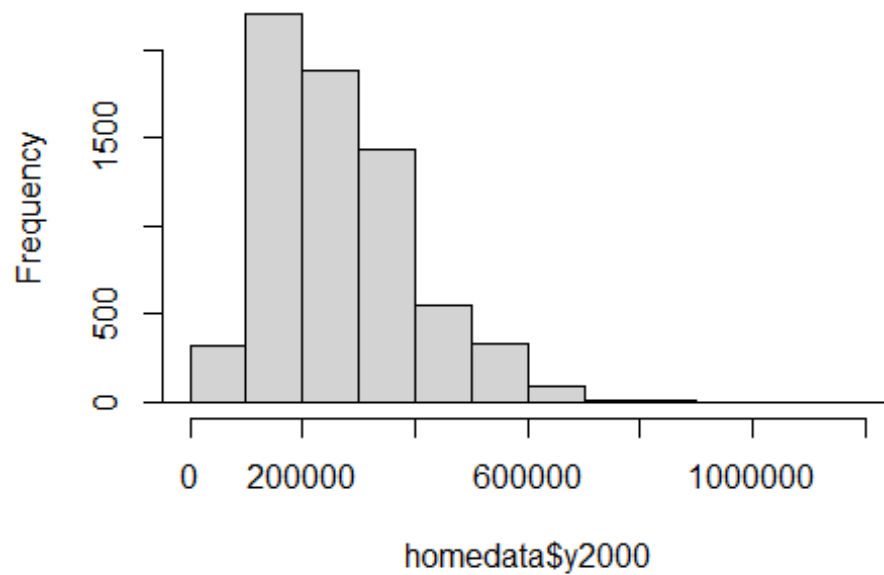
```
#3.  
#графиките за къщите от 1970 и 2000г.  
hist(homedata$y1970)  
lines(density(homedata$y1970, na.rm = T))
```

Histogram of homedata\$y1970

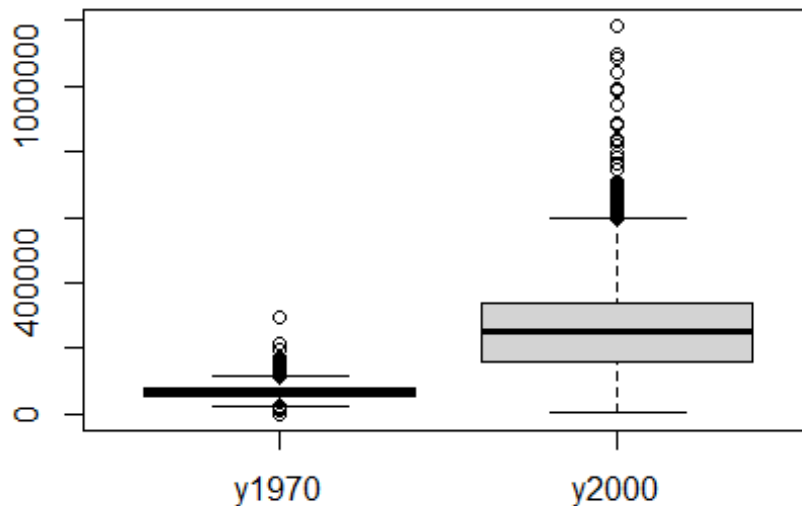


```
hist(homedata$y2000)  
lines(density(homedata$y2000, na.rm = T))
```

Histogram of homedata\$y2000



```
#сравняваме цените на къщите през 1970 и 2000г. и тяхната корелация  
boxplot(homedata)
```



```
correlation <- cor(homedata$y1970, homedata$y2000)
```

```
#4.  
#View(anscombe)  
#boxplot(anscombe)
```

От упражнение 4

```
dice = function(N = 100){  
  samples <- sample(1:6, size = 100, replace = TRUE)  
  
  result <- sum(samples == 6)  
  
  result  
}  
  
#емпирична вероятност  
dice() / 100  
  
## [1] 0.15  
  
birthdays = function(p = 0.5){
```

```

prob = 1
for(i in 1:365){
  prob = prob * (366 - i) / 365

  if(prob < 1 - p) break
}
return(i)
}

birthdays()
## [1] 23

game_one = function(father, mother){

  wins = 0

  for(i in 1:1000){
    vs_mom <- sample(0:1, 2, replace = T, prob = c(1 - mother, mother))
    vs_dad <- sample(0:1, 1, replace = T, prob = c(1 - father, father))

    if(vs_mom[1] == 1 & vs_dad == 1 | vs_mom[2] == 1 & vs_dad == 1){
      wins = wins + 1
    }

  }
  return(wins/1000)
}

game_one(0.3, 0.4)
## [1] 0.184

#4.

presents = function(n = 20){

  for(j in 1:10000){
    counter = 1

    x <- sample(1:n, n, replace = FALSE)

    for(i in 1:20){

      if(i == x[i]){
        counter = counter + 1
        break
      }
    }
    return(n - counter)
  }
}

```

```

    }
  }

presents() / 10000

## [1] 0.0019

#5.

coins = function(){
  for(i in 1:10000){

    x <- sample(0:1, 5, replace = T)

    if(x[1] == 1 & x[2] == 1 & x[3] == 0 & x[4] == 1 & x[5] == 0){
      break
    }
  }
  return(i)
}

coins()

## [1] 88

```

От упражнение 5

```

#вероятността да се паднат по-малко от 5 шестници при хвърляне на 30 зара
rbinom(q = 4, size = 30, prob = 1/6)

## [1] 0.4243389

#взимаме извадка от 10000 по 30 хвърляния на зар и го правим на таблица -
това е емп. вер.
thrown_dices <- rbinom(n = 10000, size = 30, prob = 1/6)

thrown_dices %>% table() %>% prop.table()

## .
##      0      1      2      3      4      5      6      7      8      9
10
## 0.0045 0.0280 0.0722 0.1370 0.1852 0.1828 0.1612 0.1090 0.0672 0.0330
0.0135
##      11      12      13      16
## 0.0044 0.0014 0.0005 0.0001

#това е теоритичната вероятност
dbinom(0:6, size = 30, prob = 1/6)

## [1] 0.00421272 0.02527632 0.07330133 0.13682915 0.18471936 0.19210813
0.16009011

```

```

#с вероятност 0,75 да се паднат повече от колко шестци
#понеже нямаме ф-я за повече от ние ще променим твърдението с неговото
обратно
#понеже qbinom показва колко най-много шестци ще се паднат за някаква
вероятност
#тоест ние го променяме колко най-много ще се паднат за 0.25 вероятност
qbinom(p = 0.25, size = 30, prob = 1/6)

## [1] 4

qbinom(p = 0.75, size = 30, prob = 1/6, lower.tail = FALSE)

## [1] 4

#2.

#имаме пет неуспеха преди 3тия успех като вероятността за успех е 0.2
# x е квантил
dnbinom(x = 5, size = 3, prob = 0.2)

## [1] 0.05505024

#вероятност да са му нужни повече от 6 изстрела
#q е бр. неуспехи
pnbinom(q = 3, size = 3, prob = 0.2, lower.tail = FALSE)

## [1] 0.90112

#вероятност да му трябват между 5 и 8 изстрела вкл.
pnbinom(q = 5, size = 3, prob = 0.2) - pnbinom(1,3,0.2)

## [1] 0.1758822

#3.
balls = function(){

  total = 13
  white_balls = 7
  black_balls = 6

  num_white = 0

  for(i in 1:8){

    white_p = white_balls / total
    black_p = black_balls / total

    x <- sample(c(0,1), size = 1, prob = c(black_p, white_p))

    if(x == 0){
      black_balls = black_balls - 1
    }
  }
}

```

```

    }
    else{
      white_balls = white_balls - 1
      num_white = num_white + 1
    }

    total = total - 1
  }
  num_white
}

replicated <- replicate(1000, balls())

mean(replicated)
## [1] 4.269

sd(replicated)
## [1] 0.9140429

min(replicated)
## [1] 2

max(replicated)
## [1] 7

sum(replicated == 3) / 1000
## [1] 0.175

#емпирична вероятност
emp <- replicated %>% table() %>% prop.table()

#теоритична вероятност
theor <- dhyper(0:8, 7, 6, 8)

hist(c(emp, theor) , beside = T)

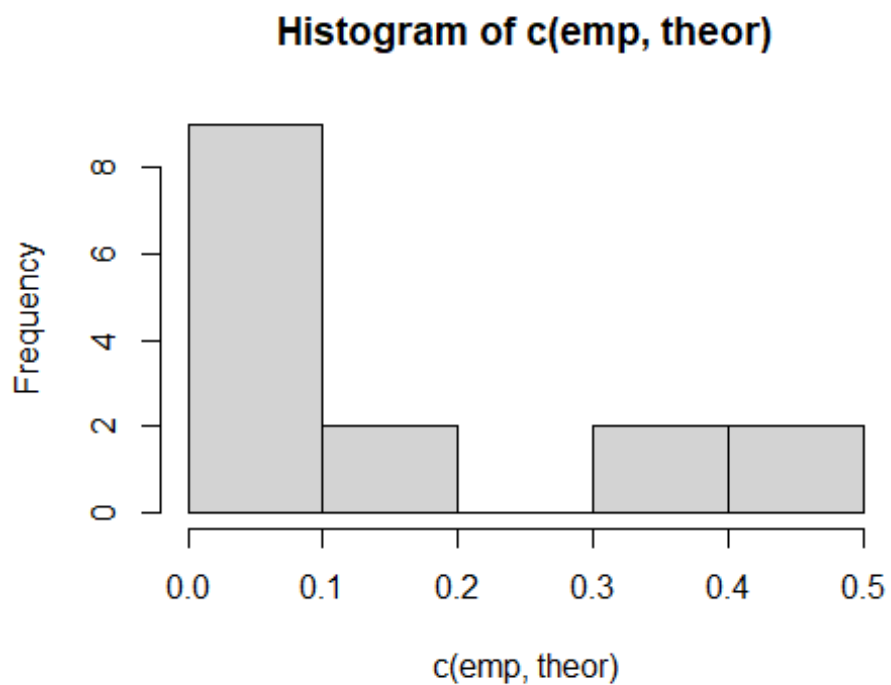
## Warning in plot.window(xlim, ylim, "", ...): "beside" is not a graphical
## parameter

## Warning in title(main = main, sub = sub, xlab = xlab, ylab = ylab, ...):
## "beside" is not a graphical parameter

## Warning in axis(1, ...): "beside" is not a graphical parameter

## Warning in axis(2, ...): "beside" is not a graphical parameter

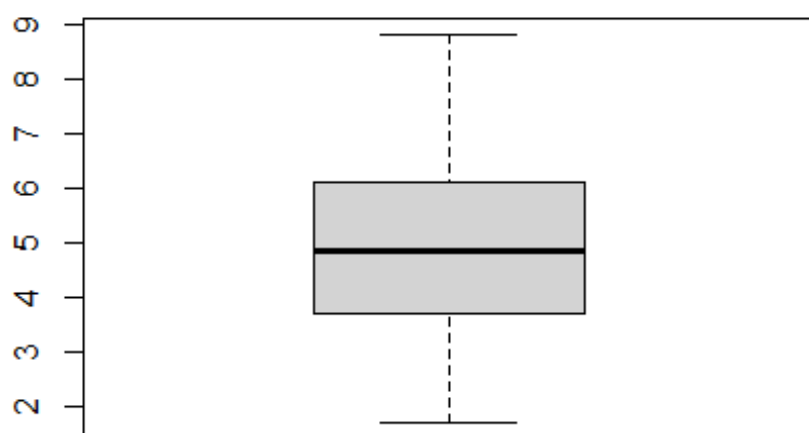
```



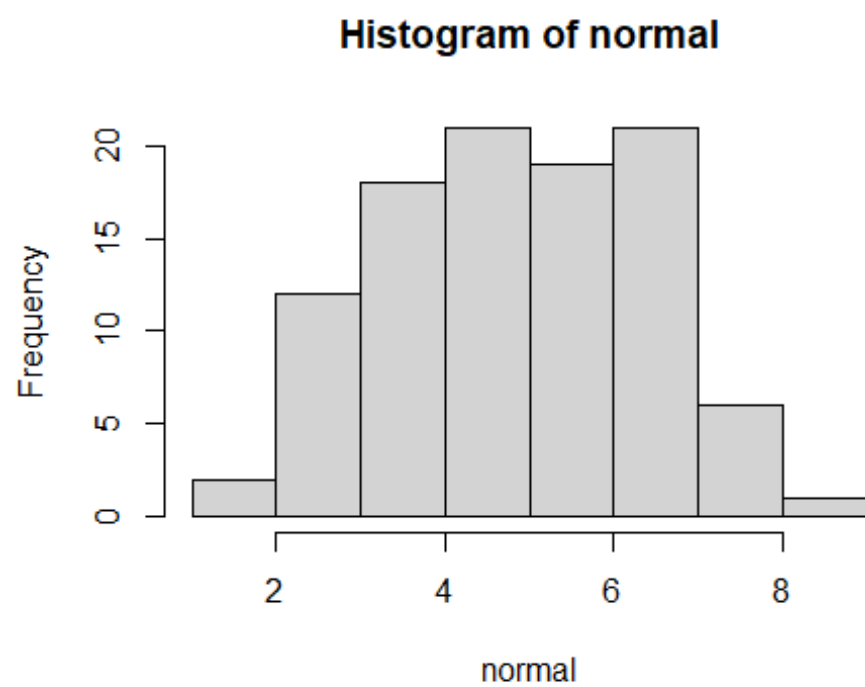
```
#4.  
n = 100  
  
dbinom(2, size = n, prob = 5/(2*n))  
## [1] 0.2587841
```

От упражнение 6

```
#нормално разпр с боксплот и хистограма  
normal <- rnorm(100, 5, sqrt(2))  
  
boxplot(normal)
```

```
hist(normal)
```



#правим някаква извадка

```
s <- seq(1, 8, 0.2)
```

#теоритична вероятност

```
dnorm(s, 5, sqrt(2))
```

```
## [1] 0.005166746 0.007631185 0.011047931 0.015677760 0.021807265  
0.029732572
```

```
## [7] 0.039735427 0.052051997 0.066836087 0.084119899 0.103776874  
0.125492144
```

```
## [13] 0.148746447 0.172818715 0.196810858 0.219695645 0.240385325  
0.257815227
```

```
## [19] 0.271033697 0.279287902 0.282094792 0.279287902 0.271033697  
0.257815227
```

```
## [25] 0.240385325 0.219695645 0.196810858 0.172818715 0.148746447  
0.125492144
```

```
## [31] 0.103776874 0.084119899 0.066836087 0.052051997 0.039735427  
0.029732572
```

#2.

#n - брой сл. в

#k - брой стойности, които ни дава всяко разпределение

fn - distribution function and ... is her arguments

```
xsim <- function(n, k, fn, ...){
```

#вектор пълен с нули. В него се събират стойностите поиндексно на всяко разпределение

```
s <- rep(0, k)
```

```
for (i in 1:n) {
```

```
  s <- s + fn(k, ... )
```

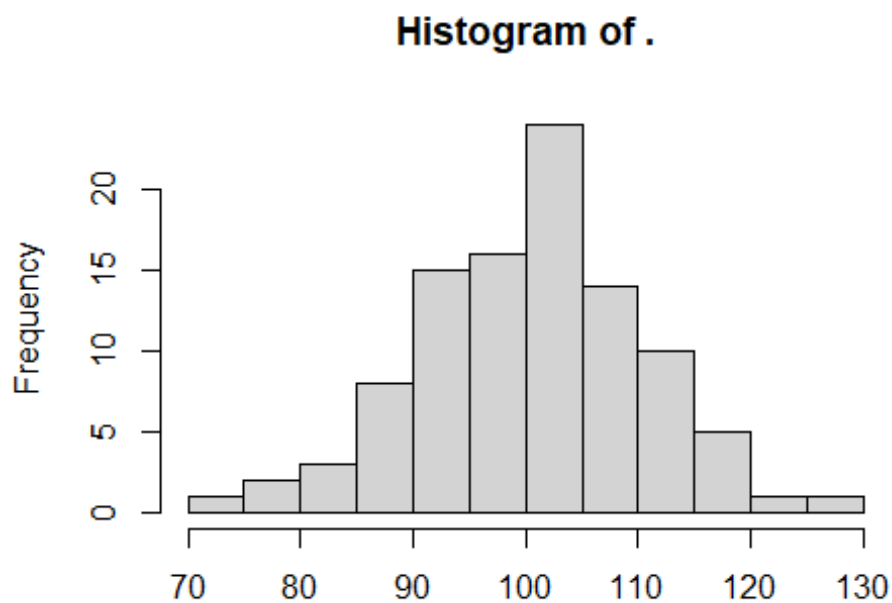
```
}
```

#върща се вектор със сумата поиндексно на всички разпределения

```
s
```

```
}
```

```
xsim(100, 100, rexp) %>% hist()
```



#това ни показва граничната теор - т.е при сумиране на независими еднакво разпр сл.в

#че се получава нормално разпределение

#4.

#пънеша по-малки от 20 т.е трето качество

```
small <- pnorm(20, 25, 6)
```

#първата половина от по-големите

```
medium <- (1 - small) / 2
```

```
big <- medium
```

#колко да е голям за да бъде трето качество

```
qnorm(big + medium, 25, 6)
```

```
## [1] 30
```

От упражнение 7

#1. а)ако ни е известно станд отклонение

```
n <- 20
```

```
sd <- 2
```

```
x <- rnorm(n, 3, sd)
```

```
q <- qnorm(0.975, 3, 2)
```

```

left_interval <- mean(x) - q * (sd / sqrt(n))
right_interval <- mean(x) + q * (sd / sqrt(n))

#b)ако не ни е известно станд отклонение

n <- 20

theor_mean <- 3
theor_sd <- 2

x <- rnorm(n, theor_mean, theor_sd)

m <- mean(x)

sd <- sd(x)

q <- qt(p = 0.975, df = n - 1)

left_interval <- m - q * sd / sqrt(n)
right_interval <- m + q * sd / sqrt(n)

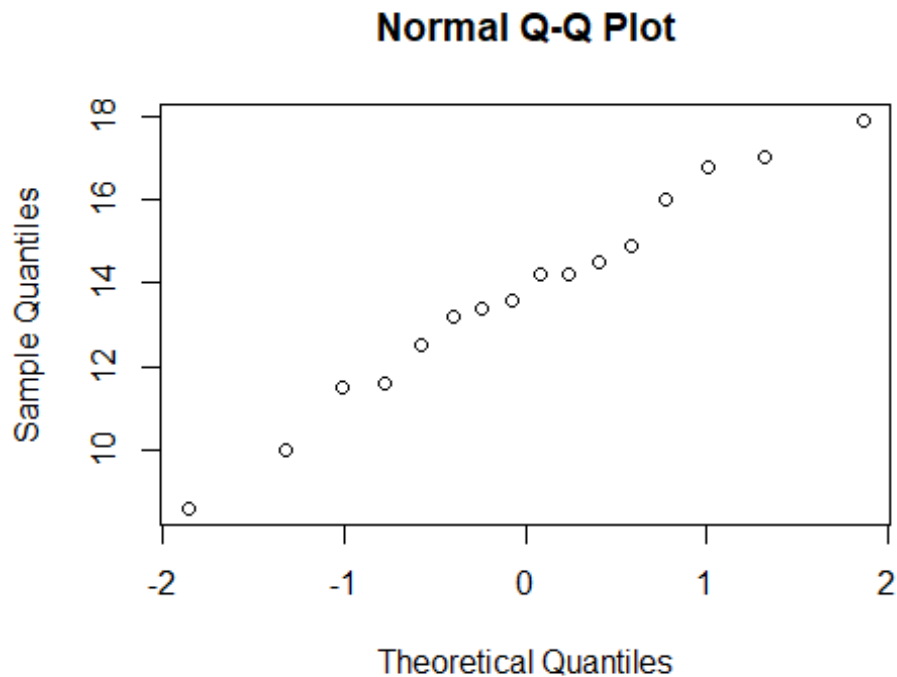
#можем и така да намерим доверителния интервал ако знаем че x е нормално
#разпределено
t.test(x)

##
## One Sample t-test
##
## data: x
## t = 6.8338, df = 19, p-value = 1.6e-06
## alternative hypothesis: true mean is not equal to 0
## 95 percent confidence interval:
##  2.276277 4.286187
## sample estimates:
## mean of x
##  3.281232

#2.
data1 <- c(10.0, 13.6, 13.2, 11.6, 12.5, 14.2, 14.9, 14.5, 13.4, 8.6, 11.5,
16.0, 14.2, 16.8, 17.9, 17.0)

#проверяваме дали е нормално разпределена
qqnorm(data1)

```



#правим доверителните интервали

```
t.test(data1)
```

```
##
```

```
## One Sample t-test
```

```
##
```

```
## data: data1
```

```
## t = 21.65, df = 15, p-value = 9.976e-13
```

```
## alternative hypothesis: true mean is not equal to 0
```

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

```
## 12.39066 15.09684
```

```
## sample estimates:
```

```
## mean of x
```

```
## 13.74375
```

```
t.test(data1, conf.level = 0.90)
```

```
##
```

```
## One Sample t-test
```

```
##
```

```
## data: data1
```

```
## t = 21.65, df = 15, p-value = 9.976e-13
```

```
## alternative hypothesis: true mean is not equal to 0
```

```
## 90 percent confidence interval:
```

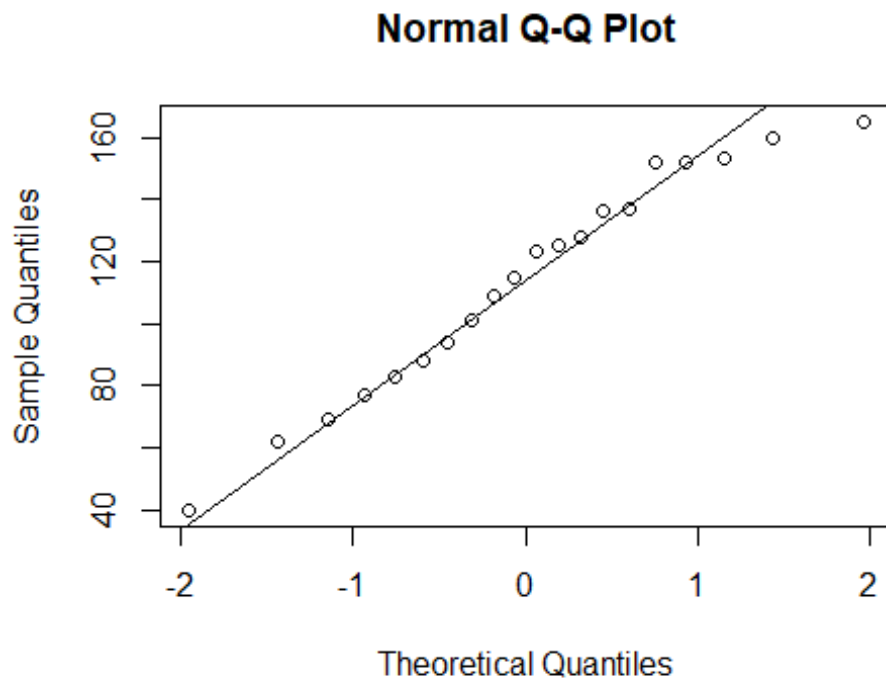
```
## 12.63088 14.85662
```

```
## sample estimates:
```

```
## mean of x
## 13.74375
```

#3.a)

```
qqnorm(rat)
qqline(rat)
```



```
t.test(rat, conf.level = 0.96)
```

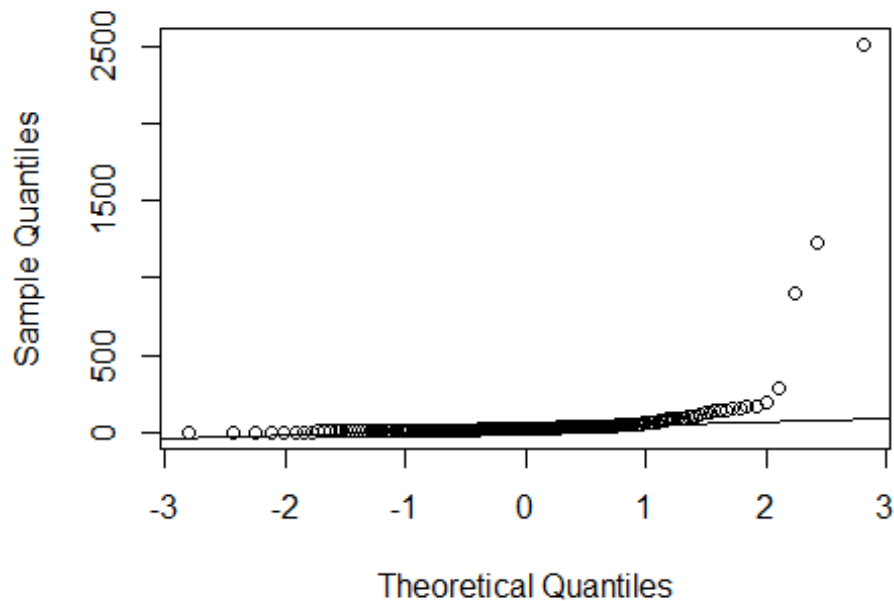
```
##
## One Sample t-test
##
## data: rat
## t = 14.176, df = 19, p-value = 1.48e-11
## alternative hypothesis: true mean is not equal to 0
## 96 percent confidence interval:
## 95.80624 131.09376
## sample estimates:
## mean of x
## 113.45
```

#b)

#данните не са нормално разпределени затова ползваме wilcox.test

```
qqnorm(exec.pay)
qqline(exec.pay)
```

Normal Q-Q Plot



```
wilcox.test(exec.pay, conf.int = T, conf.level = 0.96)

##
## Wilcoxon signed rank test with continuity correction
##
## data:  exec.pay
## V = 19306, p-value < 2.2e-16
## alternative hypothesis: true location is not equal to 0
## 96 percent confidence interval:
##  25.99996 33.00003
## sample estimates:
## (pseudo)median
##      29.00002

#v)

#4.когато имаме пропорции ползваме това
prop.test(87, 150, conf.level = 0.92)

##
## 1-sample proportions test with continuity correction
##
## data:  87 out of 150, null probability 0.5
## X-squared = 3.5267, df = 1, p-value = 0.06039
## alternative hypothesis: true p is not equal to 0.5
## 92 percent confidence interval:
##  0.5051991 0.6514474
```

```
## sample estimates:
##      p
## 0.58

#6.
smoke_men <- nrow(survey[survey$Sex == 'Male' & survey$Smoke == 'Never', ])

all_men <- nrow(survey[survey$Sex == 'Male', ])

prop.test(smoke_men, all_men, conf.level = 0.90)

##
## 1-sample proportions test with continuity correction
##
## data:  smoke_men out of all_men, null probability 0.5
## X-squared = 32.303, df = 1, p-value = 1.319e-08
## alternative hypothesis: true p is not equal to 0.5
## 90 percent confidence interval:
##  0.6908187 0.8260629
## sample estimates:
##           p
## 0.7647059
```

От упражнение 8

```
#1. check for norm distr
#  $H_0 - EX = 3$ 
data2 <- rnorm(100, mean = 2, sd = 2)

#p-value < 0.05 отхвърляме хипотезата
t.test(data2, mu = 3, alternative = 'two.sided')

##
## One Sample t-test
##
## data:  data2
## t = -4.4363, df = 99, p-value = 2.376e-05
## alternative hypothesis: true mean is not equal to 3
## 95 percent confidence interval:
##  1.646252 2.482987
## sample estimates:
## mean of x
##  2.06462

t.test(data2, mu = 5, alternative = 'two.sided')

##
## One Sample t-test
##
## data:  data2
## t = -13.922, df = 99, p-value < 2.2e-16
```



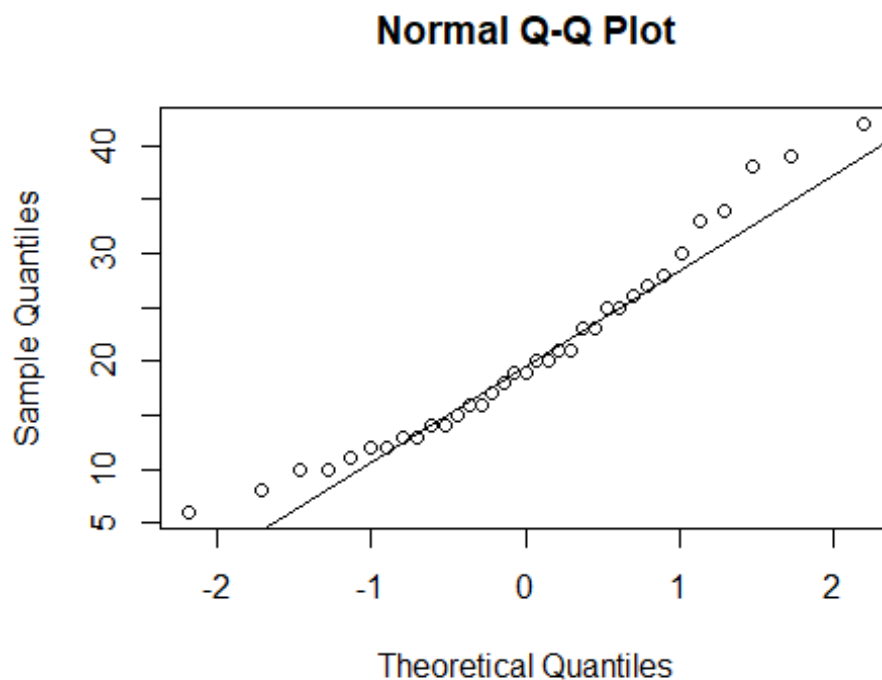
```
## alternative hypothesis: true mean is not equal to 5
## 95 percent confidence interval:
##  1.646252 2.482987
## sample estimates:
## mean of x
##  2.06462
```

```
#2.H0: дните за почивка да са 24 при n-жалуе > 0.2
data3 <- vacation
```

```
#seems normal distributed
```

```
qqnorm(data3)
```

```
qqline(data3)
```



```
#p-value < 0,2 отхвърляме нулевата хипотеза
```

```
t.test(vacation, mu = 24, alternative = 'two.sided')
```

```
##
```

```
## One Sample t-test
```

```
##
```

```
## data: vacation
```

```
## t = -2.2584, df = 34, p-value = 0.03045
```

```
## alternative hypothesis: true mean is not equal to 24
```

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

```
## 17.37768 23.65089
```

```
## sample estimates:
```

```
## mean of x
## 20.51429

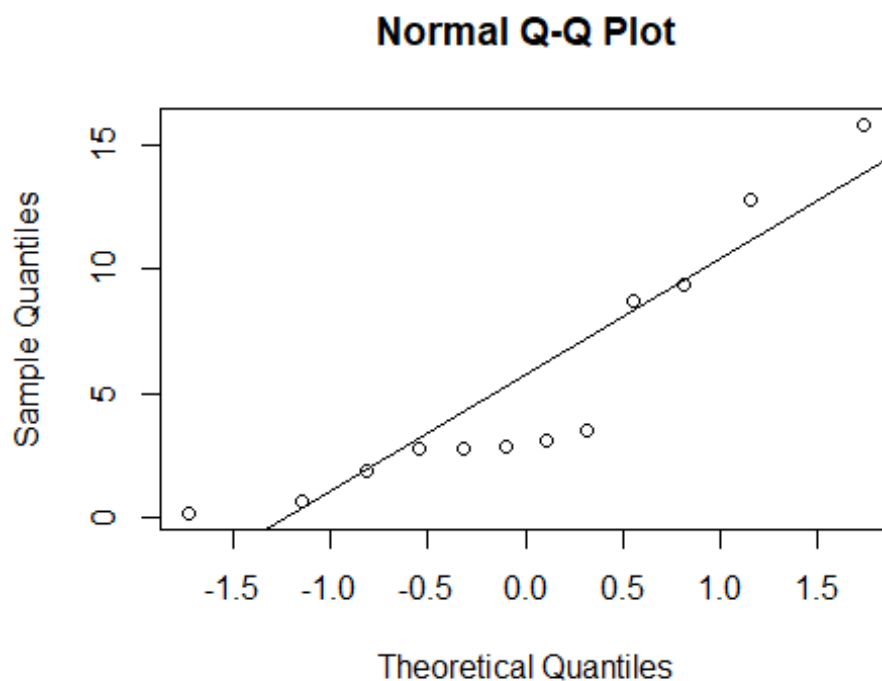
#3.H0 - 50 процента са доволни
# H1 - < 50 процента са доволни

prop.test(42, 100, p = 0.5, alternative = 'less')

##
## 1-sample proportions test with continuity correction
##
## data: 42 out of 100, null probability 0.5
## X-squared = 2.25, df = 1, p-value = 0.06681
## alternative hypothesis: true p is less than 0.5
## 95 percent confidence interval:
## 0.0000000 0.5072341
## sample estimates:
## p
## 0.42

#4.H0 - 5 minutes on the phone H1 - more than 5 mins
data4 <- c(12.8, 3.5, 2.9, 9.4, 8.7, 0.7, 0.2, 2.8, 1.9, 2.8, 3.1, 15.8)

qqnorm(data4)
qqline(data4)
```



```

#not normal distribution
shapiro.test(data4)

##
##  Shapiro-Wilk normality test
##
## data:  data4
## W = 0.83988, p-value = 0.0276

#we accept h0
wilcox.test(data4, mu = 5, alternative = 'greater')

## Warning in wilcox.test.default(data4, mu = 5, alternative = "greater"):
## cannot
## compute exact p-value with ties

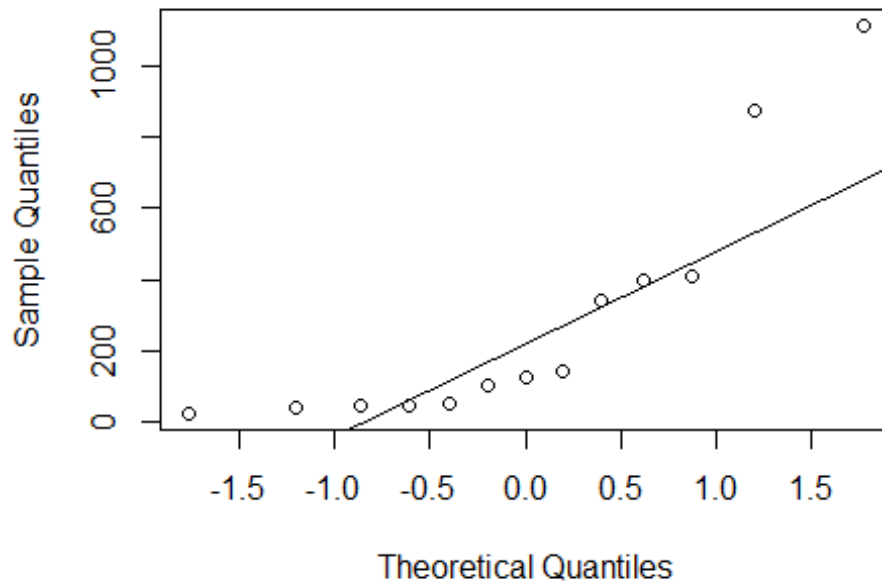
##
##  Wilcoxon signed rank test with continuity correction
##
## data:  data4
## V = 39, p-value = 0.5156
## alternative hypothesis: true location is greater than 5

#5.h0 - Live more than 100 days h1- Less than 100 days
data5 <- cancer$stomach

qqnorm(data5)
qqline(data5)

```

Normal Q-Q Plot



```
#not normal distr
shapiro.test(data5)

##
##  Shapiro-Wilk normality test
##
## data:  data5
## W = 0.75473, p-value = 0.002075

#we accept h0
wilcox.test(data5, mu = 100, alternative = 'less')

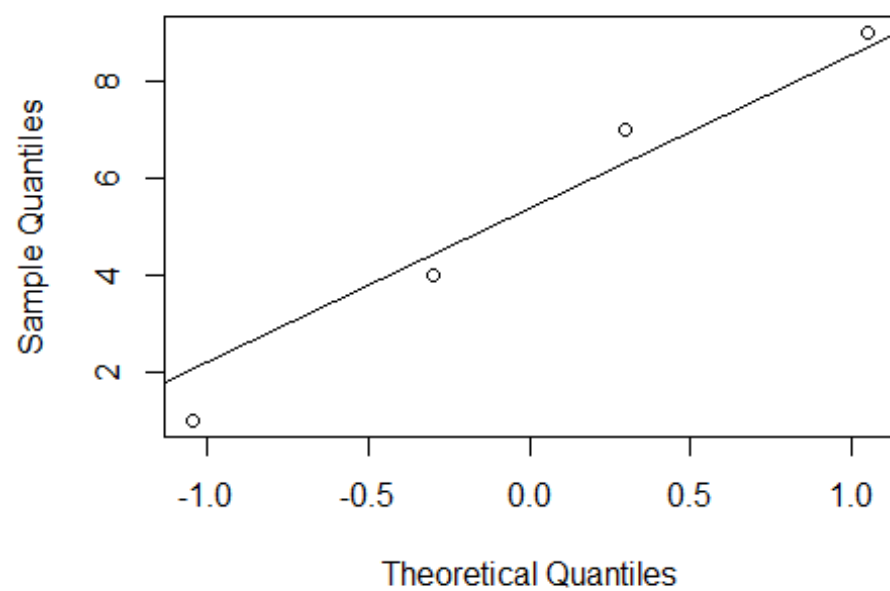
##
##  Wilcoxon signed rank exact test
##
## data:  data5
## V = 61, p-value = 0.8633
## alternative hypothesis: true location is less than 100
```

От упражнение 9

```
#1. H0 - equal h1 - not equal
x <- c(4, 1, 7, 9)
y <- c(10, 3, 2, 11)

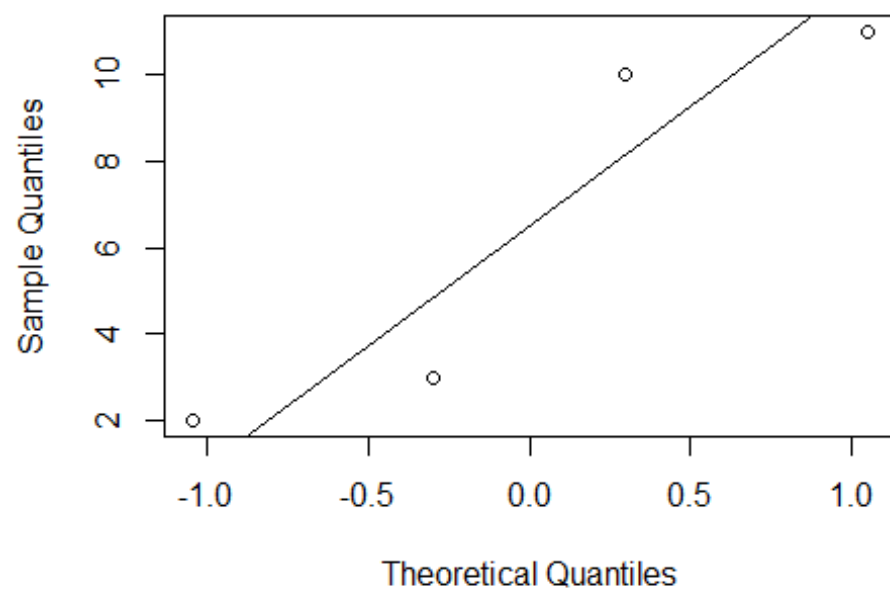
#не са норм разпр
qqnorm(x)
qqline(x)
```

Normal Q-Q Plot



```
qqnorm(y)  
qqline(y)
```

Normal Q-Q Plot



```

#приемаме нулевата хипотеза
wilcox.test(x, y, alternative = 'two.sided')

##
## Wilcoxon rank sum exact test
##
## data: x and y
## W = 6, p-value = 0.6857
## alternative hypothesis: true location shift is not equal to 0

#2.H0 - equal H1 - greater

#we accept h1
prop.test(c(351, 71), c(605, 195), alternative = 'greater')

##
## 2-sample test for equality of proportions with continuity correction
##
## data: c(351, 71) out of c(605, 195)
## X-squared = 26.761, df = 1, p-value = 1.151e-07
## alternative hypothesis: greater
## 95 percent confidence interval:
## 0.1470851 1.0000000
## sample estimates:
## prop 1 prop 2
## 0.5801653 0.3641026

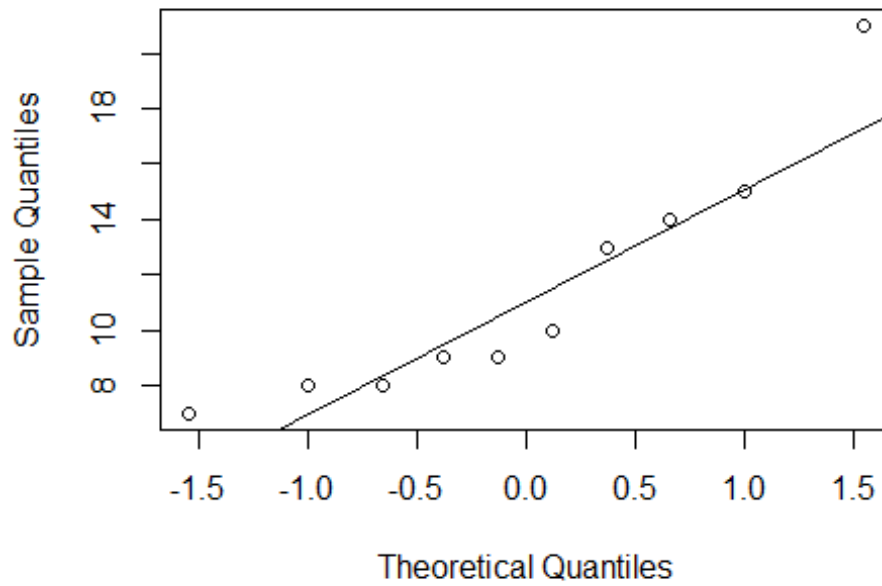
#3.H0 - less H1 - more

before <- c( 15, 10, 13, 7, 9, 8, 21, 9, 14, 8)
after <- c(15, 14, 12, 8, 14, 10, 7, 16, 10, 15, 12)

#normal distr
qqnorm(before)
qqline(before)

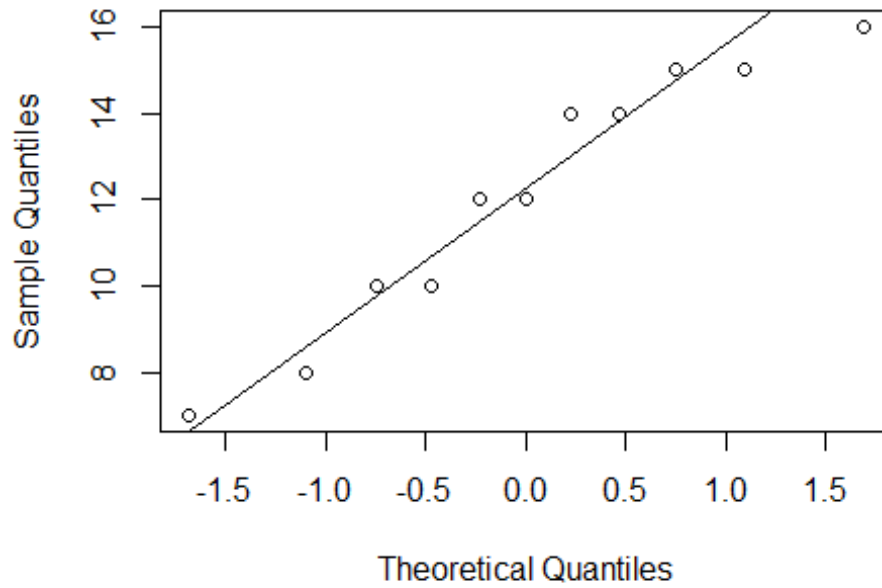
```

Normal Q-Q Plot



```
qqnorm(after)  
qqline(after)
```

Normal Q-Q Plot



```

t.test(before, after, alternative = 'greater')

##
##  Welch Two Sample t-test
##
## data:  before and after
## t = -0.41894, df = 15.853, p-value = 0.6596
## alternative hypothesis: true difference in means is greater than 0
## 95 percent confidence interval:
##  -3.571812      Inf
## sample estimates:
## mean of x mean of y
##  11.40000  12.09091

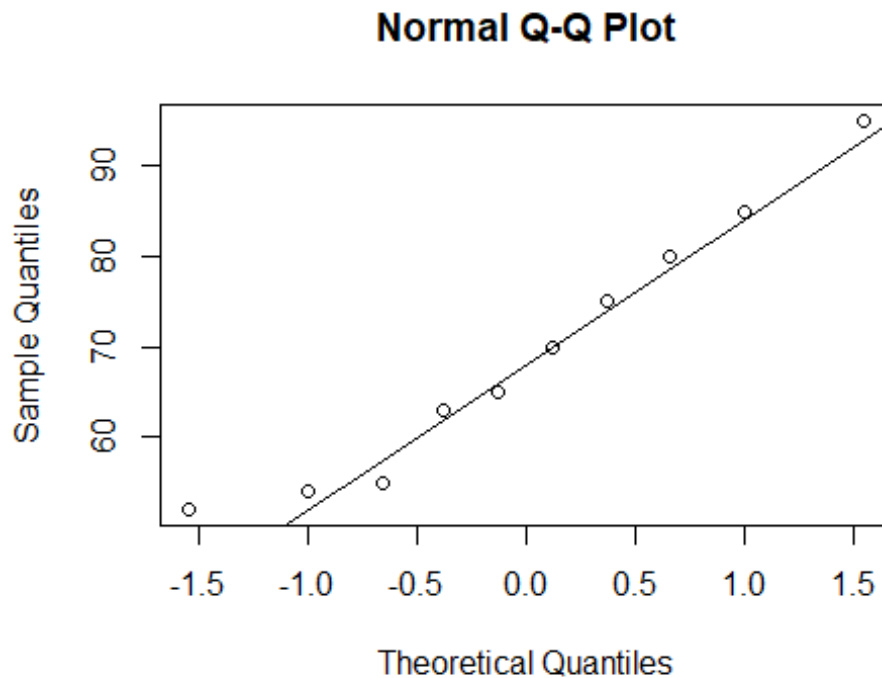
#4.X0 - че са равни X1 - че са различни

radar1 <- c(70, 85, 63, 54, 65, 80, 75, 95, 52, 55)

radar2 <- c(72, 86, 62, 55, 63, 80, 78, 90, 53, 57)

#they are normally distributed
qqnorm(radar1)
qqline(radar1)

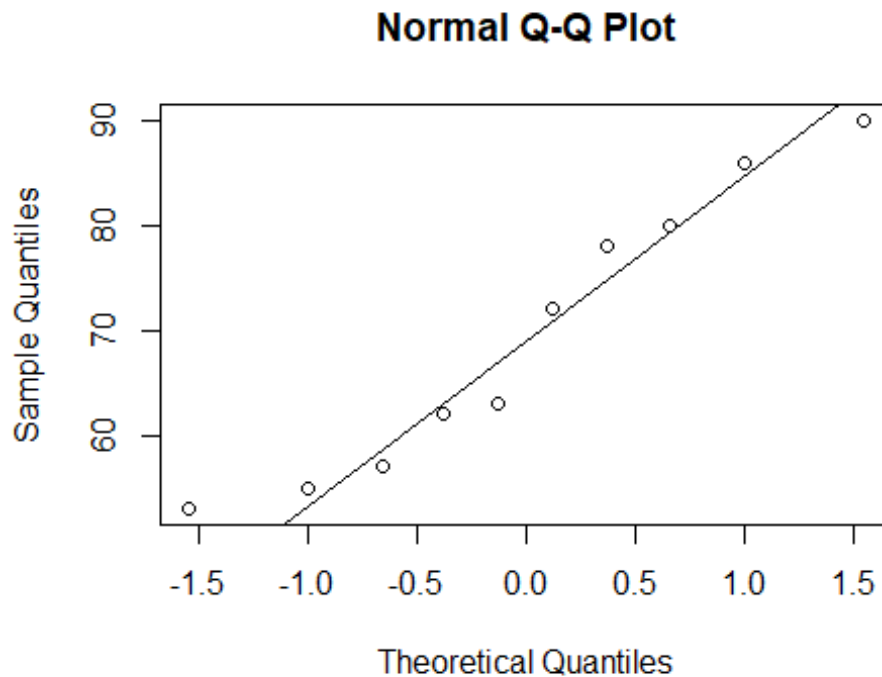
```



```

qqnorm(radar2)
qqline(radar2)

```

```
t.test(radar1, radar2, alternative = 'two.sided', paired = T)

##
## Paired t-test
##
## data: radar1 and radar2
## t = -0.26941, df = 9, p-value = 0.7937
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -1.879354 1.479354
## sample estimates:
## mean of the differences
## -0.2
```

От упражнение 10

```
#1.H0 - data elements are equal H1- they are not

data6 <- c(125, 410, 310, 300, 318, 298, 148)

#we cant accept H0
chisq.test(data6)

##
## Chi-squared test for given probabilities
##
```

```

## data: data6
## X-squared = 223.84, df = 6, p-value < 2.2e-16

#2.H0 - all digits have the same prob H1 - they don't

#взимаме първите 200 цифри и виждаме колко често всяка една се среща
data7 <- pi2000[1:200] %>% table()

#правим хи-квадрат теста
#можем да приемем нулевата хипотеза
chisq.test(data7)

##
## Chi-squared test for given probabilities
##
## data: data7
## X-squared = 7.2, df = 9, p-value = 0.6163

#3.
#теоритичните вероятности за срещането на буквите в англ език
probs <- c(0.1270, 0.0956, 0.0817, 0.0751, 0.0697, 0.0675, 0.4834)

#срещането на буквите от нашия текст
letters <- c(102, 108, 90, 95, 82, 40, 519)

#правим проверка дали са равни
chisq.test(letters, p = probs)

##
## Chi-squared test for given probabilities
##
## data: letters
## X-squared = 26.396, df = 6, p-value = 0.0001878

#p-value е много малко затова отхвърляме хипотезата

#4.искаме да видим дали колан / без колан са независими сл.б Това става като
подадем
#на ф-ята матрица и тя прави проверката, т.е нулевата хипотеза е, че данните
са независими
# иначе са зависими

belt <- c(12813, 647, 359, 42)

nobelt <- c(65963, 4000, 2642, 303)

data8 <- matrix(belt, nrow = 1, ncol = 4)

data9 <- rbind(data8, nobelt)

```

#p-value е много малко следователно отхвърляме нулевата хипотеза, т.е са зависими

#колан влияе на нараняването при катастрофа

```
chisq.test(data9)
```

```
##
```

```
## Pearson's Chi-squared test
```

```
##
```

```
## data: data9
```

```
## X-squared = 59.224, df = 3, p-value = 8.61e-13
```

#5.

```
mon <- c(44, 14, 15, 3)
```

```
tues <- c(74, 25, 20, 5)
```

```
wed <- c(79, 27, 20, 5)
```

```
thurs <- c(72, 24, 23, 0)
```

```
fri <- c(31, 10, 9, 0)
```

```
final <- matrix(c(mon, tues, wed, thurs, fri), nrow = 5)
```

#отхвърляме нулевата хипотеза че са независими , т.е има връзка между деня и качеството на стоката

```
chisq.test(final)
```

```
##
```

```
## Pearson's Chi-squared test
```

```
##
```

```
## data: final
```

```
## X-squared = 350.71, df = 12, p-value < 2.2e-16
```

От упражнение 11

#1.Създаваме дата фрейм за данните

```
patients_df <- data.frame(
```

```
  age = c(18, 23, 25, 35, 65, 54, 34, 56, 72, 19, 23, 42, 18, 39, 37),
```

```
  max_pulse = c(202, 186, 187, 180, 156, 169, 174, 172, 153, 199, 193, 174, 198, 183, 178)
```

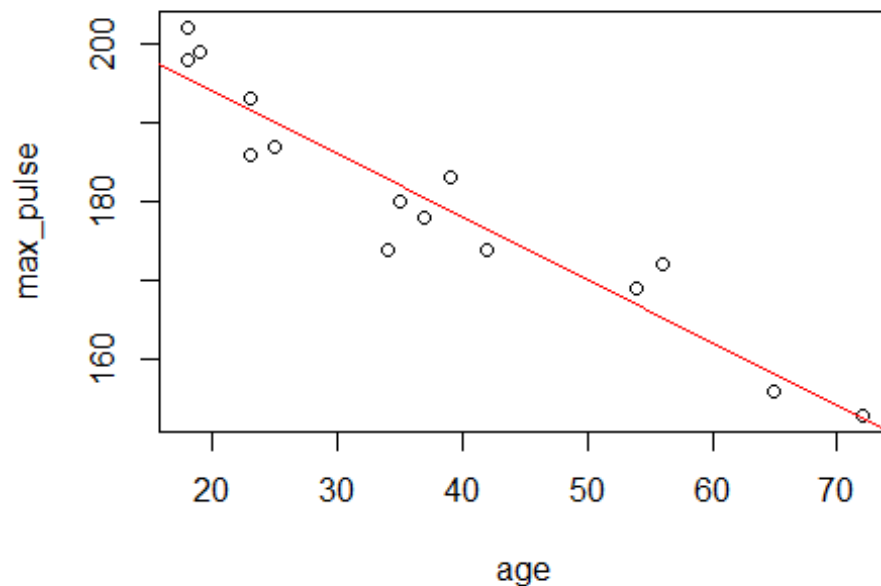
```
)
```

#това е за модел на линейната регресия

```
model1 <- lm(patients_df$max_pulse ~ patients_df$age, data = patients_df)
```

```
plot(patients_df)
```

```
abline(model1, col = "red")
```



```
summ_lm <- summary(model1)

n <- nrow(patients_df)

# Тестване на хипотезата, че бета1 = -1
# H0 :- "бета_1 = -1"

# Стандартно отклонение(грешка) на оценката за бета1
std_b1 <- summ_lm$coefficients[2, 2]

# оценката за бета1
est_b1 <- summ_lm$coefficients[2, 1]

# Параметър за бета1 под нулева хипотеза
b1_null_hyp <- -1

# Изграждане на t-статистика
t_statistic <- (est_b1 - b1_null_hyp) / std_b1

# Вероятност да наблюдаваме тази t-статистика (или по-крайна) при положение,
# че е вярна нулевата хипотеза
pval <- 2 * pt(t_statistic, n - 2, lower.tail = FALSE)

# Прогнозиране за възрасти 30, 40 и 50
```

```

predict.lm(
  model1,
  newdata = data.frame(age = c(30, 40, 50)),
  interval = "confidence",
  level = 0.9
)

## Warning: 'newdata' had 3 rows but variables found have 15 rows

##           fit      lwr      upr
## 1  195.6894 192.5083 198.8705
## 2  191.7007 188.9557 194.4458
## 3  190.1053 187.5137 192.6969
## 4  182.1280 180.0149 184.2411
## 5  158.1962 154.1798 162.2127
## 6  166.9712 164.0309 169.9116
## 7  182.9258 180.7922 185.0593
## 8  165.3758 162.2564 168.4952
## 9  152.6121 147.8341 157.3902
## 10 194.8917 191.8028 197.9805
## 11 191.7007 188.9557 194.4458
## 12 176.5439 174.3723 178.7155
## 13 195.6894 192.5083 198.8705
## 14 178.9371 176.8337 181.0405
## 15 180.5326 178.4390 182.6262

```

От упражнение 12

От упражнение 13

```

#нулевата хипотеза е че имат равни средни трите извадки от данни
#правим дата фрейм с данните от задачата
exams_df <- data.frame(
  examiner1 = c(5, 4, 4, 6, 4, 6, 3, 3, 4, 5),
  examiner2 = c(3, 2, 4, 5, 3, 4, 3, 4, 2, 4),
  examiner3 = c(4, 6, 4, 2, 4, 5, 5, 3, 6, 4)
)

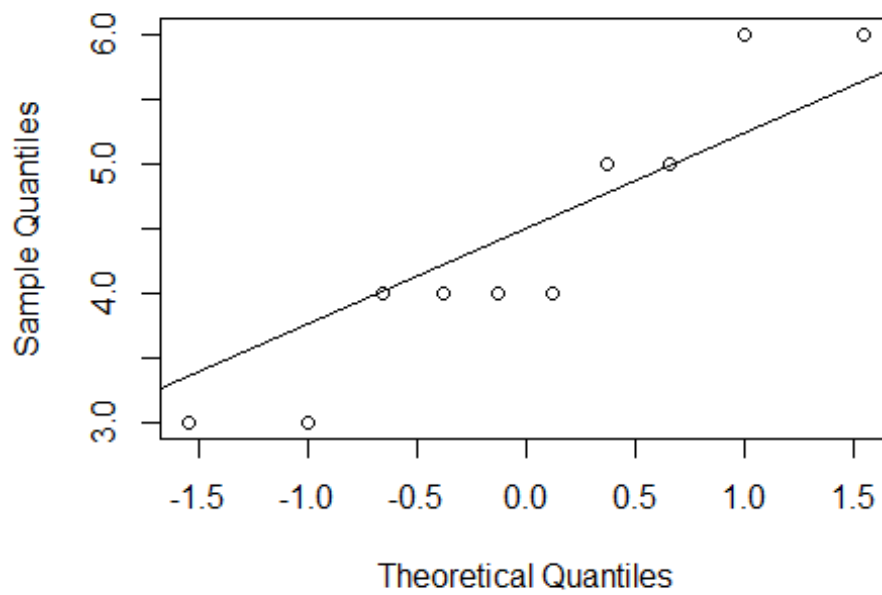
stacked_exam_df <- stack(exams_df)

#гледаме дали са нормално разпределени данните

qqnorm(exams_df$examiner1)
qqline(exams_df$examiner1)

```

Normal Q-Q Plot



```
shapiro.test(exams_df$examinor1)
```

```
##
```

```
##  Shapiro-Wilk normality test
```

```
##
```

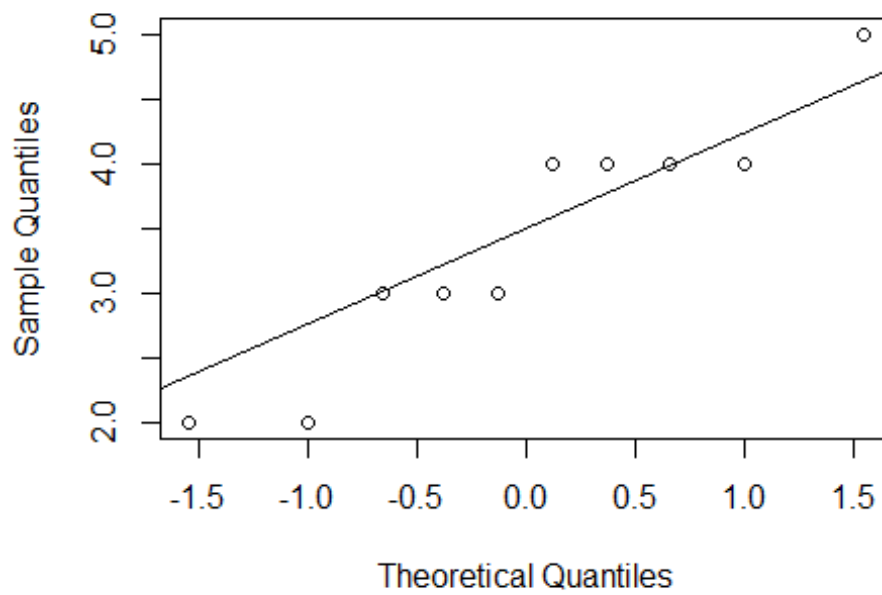
```
## data:  exams_df$examinor1
```

```
## W = 0.89165, p-value = 0.177
```

```
qqnorm(exams_df$examinor2)
```

```
qqline(exams_df$examinor2)
```

Normal Q-Q Plot

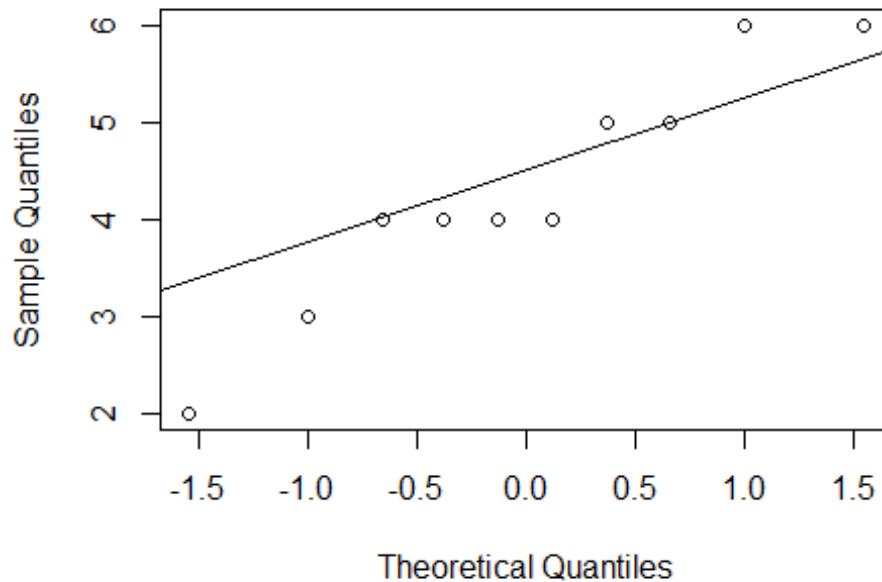


```
shapiro.test(exams_df$examinor2)

##
##  Shapiro-Wilk normality test
##
## data:  exams_df$examinor2
## W = 0.90444, p-value = 0.2449

qqnorm(exams_df$examinor3)
qqline(exams_df$examinor3)
```

Normal Q-Q Plot



```
shapiro.test(exams_df$examinor3)
```

```
##  
## Shapiro-Wilk normality test  
##  
## data: exams_df$examinor3  
## W = 0.92883, p-value = 0.4365
```

*#и трите са нормално разпределени тоест можем да направим тест дали имат
еднакво средно ако са норм разпр*

```
oneway.test(values ~ ind, data = stacked_exam_df)
```

```
##  
## One-way analysis of means (not assuming equal variances)  
##  
## data: values and ind  
## F = 2.7825, num df = 2.000, denom df = 17.811, p-value = 0.0888
```

*#Не можем да отхвърлим хипотезата че имат равни средни
#друг начин да се провери същата хипотеза*

```
anova(lm(values ~ ind, data = stacked_exam_df))
```

```
## Analysis of Variance Table  
##  
## Response: values
```



```
##           Df Sum Sq Mean Sq F value Pr(>F)
## ind       2  6.067   3.0333   2.4894 0.1018
## Residuals 27 32.900   1.2185
```

#2.

```
groupC <- InsectSprays$count[InsectSprays$spray == 'C']
```

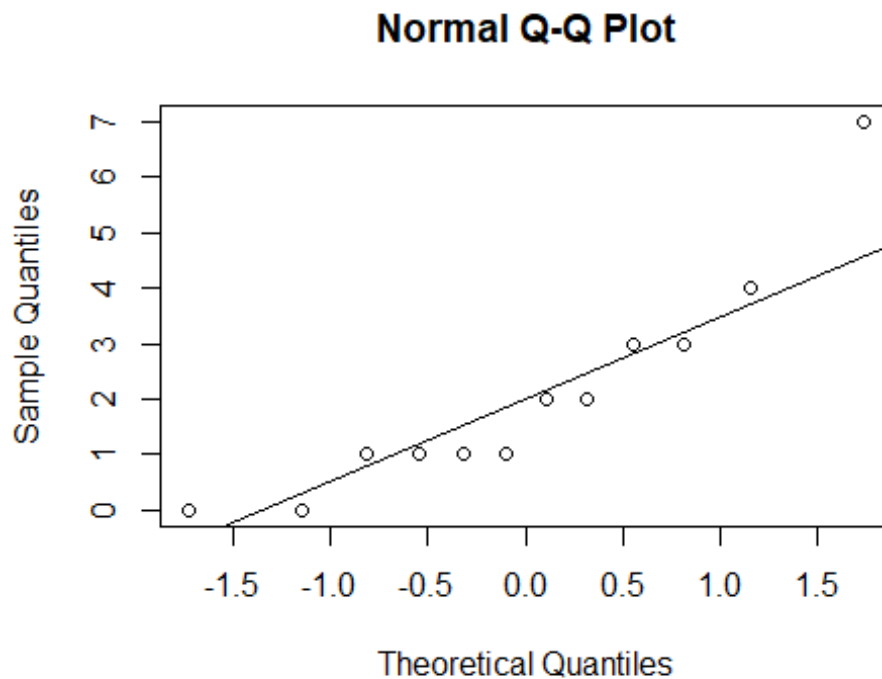
```
groupD <- InsectSprays$count[InsectSprays$spray == 'D']
```

```
groupE <- InsectSprays$count[InsectSprays$spray == 'E']
```

#изглежда ми сравнително нормално разпределени

```
qqnorm(groupC)
```

```
qqline(groupC)
```



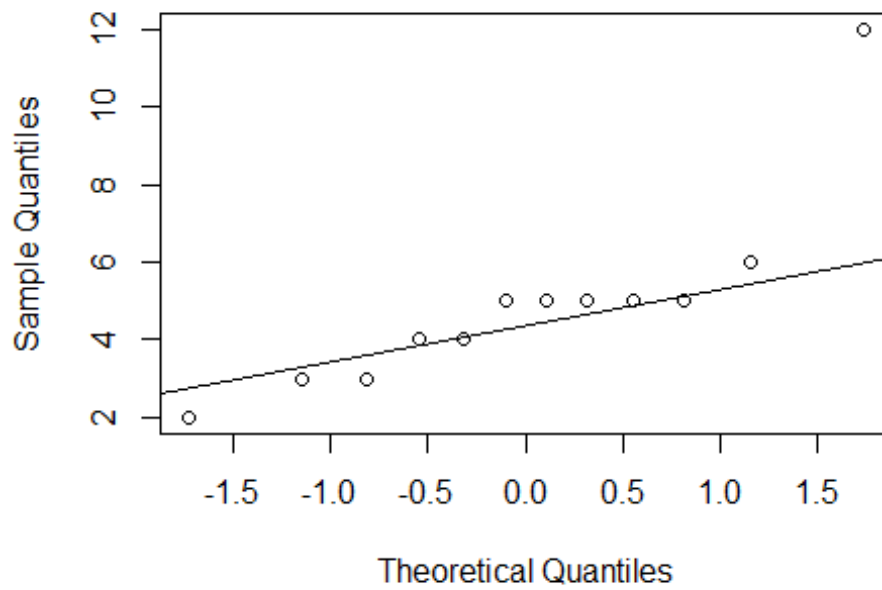
```
shapiro.test(groupC)
```

```
##
##  Shapiro-Wilk normality test
##
## data:  groupC
## W = 0.85907, p-value = 0.04759
```

```
qqnorm(groupD)
```

```
qqline(groupD)
```

Normal Q-Q Plot

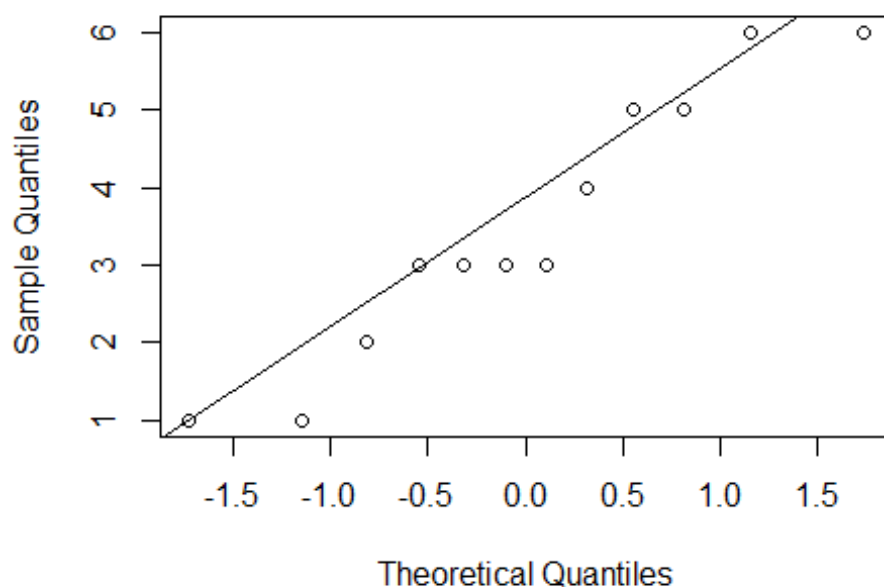


```
shapiro.test(groupD)

##
##  Shapiro-Wilk normality test
##
## data:  groupD
## W = 0.75063, p-value = 0.002713

qqnorm(groupE)
qqline(groupE)
```

Normal Q-Q Plot



```
shapiro.test(groupE)

##
##  Shapiro-Wilk normality test
##
## data:  groupE
## W = 0.92128, p-value = 0.2967

#p-value-то е много малко следователно можем да твърдим че някои от
#препаратите действат по-добре от други
oneway.test(count ~ spray, data = InsectSprays)

##
##  One-way analysis of means (not assuming equal variances)
##
## data:  count and spray
## F = 36.065, num df = 5.000, denom df = 30.043, p-value = 7.999e-12

#3.взимаме данните от файла
drug_df <- read.csv("./data.txt")

#тъй като имаме сдвоени данни, т.е даваме лекарство на един и същ пациент
#ползваме aov
aov(response ~ drug + Error(patient), data = drug_df) %>% summary()

##
## Error: patient
##           Df Sum Sq Mean Sq F value Pr(>F)
```

```
## Residuals 1 72.9 72.9
##
## Error: Within
##          Df Sum Sq Mean Sq F value Pr(>F)
## drug      1    36   36.00   0.565  0.462
## Residuals 17  1082   63.66
```

##не можем да отхвърлим хопотезата, че имаме лекарствата действат еднакво

#4.
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
## 6	5.4	3.9	1.7	0.4	setosa
## 7	4.6	3.4	1.4	0.3	setosa
## 8	5.0	3.4	1.5	0.2	setosa
## 9	4.4	2.9	1.4	0.2	setosa
## 10	4.9	3.1	1.5	0.1	setosa
## 11	5.4	3.7	1.5	0.2	setosa
## 12	4.8	3.4	1.6	0.2	setosa
## 13	4.8	3.0	1.4	0.1	setosa
## 14	4.3	3.0	1.1	0.1	setosa
## 15	5.8	4.0	1.2	0.2	setosa
## 16	5.7	4.4	1.5	0.4	setosa
## 17	5.4	3.9	1.3	0.4	setosa
## 18	5.1	3.5	1.4	0.3	setosa
## 19	5.7	3.8	1.7	0.3	setosa
## 20	5.1	3.8	1.5	0.3	setosa
## 21	5.4	3.4	1.7	0.2	setosa
## 22	5.1	3.7	1.5	0.4	setosa
## 23	4.6	3.6	1.0	0.2	setosa
## 24	5.1	3.3	1.7	0.5	setosa
## 25	4.8	3.4	1.9	0.2	setosa
## 26	5.0	3.0	1.6	0.2	setosa
## 27	5.0	3.4	1.6	0.4	setosa
## 28	5.2	3.5	1.5	0.2	setosa
## 29	5.2	3.4	1.4	0.2	setosa
## 30	4.7	3.2	1.6	0.2	setosa
## 31	4.8	3.1	1.6	0.2	setosa
## 32	5.4	3.4	1.5	0.4	setosa
## 33	5.2	4.1	1.5	0.1	setosa
## 34	5.5	4.2	1.4	0.2	setosa
## 35	4.9	3.1	1.5	0.2	setosa
## 36	5.0	3.2	1.2	0.2	setosa
## 37	5.5	3.5	1.3	0.2	setosa

## 38	4.9	3.6	1.4	0.1	setosa
## 39	4.4	3.0	1.3	0.2	setosa
## 40	5.1	3.4	1.5	0.2	setosa
## 41	5.0	3.5	1.3	0.3	setosa
## 42	4.5	2.3	1.3	0.3	setosa
## 43	4.4	3.2	1.3	0.2	setosa
## 44	5.0	3.5	1.6	0.6	setosa
## 45	5.1	3.8	1.9	0.4	setosa
## 46	4.8	3.0	1.4	0.3	setosa
## 47	5.1	3.8	1.6	0.2	setosa
## 48	4.6	3.2	1.4	0.2	setosa
## 49	5.3	3.7	1.5	0.2	setosa
## 50	5.0	3.3	1.4	0.2	setosa
## 51	7.0	3.2	4.7	1.4	versicolor
## 52	6.4	3.2	4.5	1.5	versicolor
## 53	6.9	3.1	4.9	1.5	versicolor
## 54	5.5	2.3	4.0	1.3	versicolor
## 55	6.5	2.8	4.6	1.5	versicolor
## 56	5.7	2.8	4.5	1.3	versicolor
## 57	6.3	3.3	4.7	1.6	versicolor
## 58	4.9	2.4	3.3	1.0	versicolor
## 59	6.6	2.9	4.6	1.3	versicolor
## 60	5.2	2.7	3.9	1.4	versicolor
## 61	5.0	2.0	3.5	1.0	versicolor
## 62	5.9	3.0	4.2	1.5	versicolor
## 63	6.0	2.2	4.0	1.0	versicolor
## 64	6.1	2.9	4.7	1.4	versicolor
## 65	5.6	2.9	3.6	1.3	versicolor
## 66	6.7	3.1	4.4	1.4	versicolor
## 67	5.6	3.0	4.5	1.5	versicolor
## 68	5.8	2.7	4.1	1.0	versicolor
## 69	6.2	2.2	4.5	1.5	versicolor
## 70	5.6	2.5	3.9	1.1	versicolor
## 71	5.9	3.2	4.8	1.8	versicolor
## 72	6.1	2.8	4.0	1.3	versicolor
## 73	6.3	2.5	4.9	1.5	versicolor
## 74	6.1	2.8	4.7	1.2	versicolor
## 75	6.4	2.9	4.3	1.3	versicolor
## 76	6.6	3.0	4.4	1.4	versicolor
## 77	6.8	2.8	4.8	1.4	versicolor
## 78	6.7	3.0	5.0	1.7	versicolor
## 79	6.0	2.9	4.5	1.5	versicolor
## 80	5.7	2.6	3.5	1.0	versicolor
## 81	5.5	2.4	3.8	1.1	versicolor
## 82	5.5	2.4	3.7	1.0	versicolor
## 83	5.8	2.7	3.9	1.2	versicolor
## 84	6.0	2.7	5.1	1.6	versicolor
## 85	5.4	3.0	4.5	1.5	versicolor
## 86	6.0	3.4	4.5	1.6	versicolor
## 87	6.7	3.1	4.7	1.5	versicolor

## 88	6.3	2.3	4.4	1.3 versicolor
## 89	5.6	3.0	4.1	1.3 versicolor
## 90	5.5	2.5	4.0	1.3 versicolor
## 91	5.5	2.6	4.4	1.2 versicolor
## 92	6.1	3.0	4.6	1.4 versicolor
## 93	5.8	2.6	4.0	1.2 versicolor
## 94	5.0	2.3	3.3	1.0 versicolor
## 95	5.6	2.7	4.2	1.3 versicolor
## 96	5.7	3.0	4.2	1.2 versicolor
## 97	5.7	2.9	4.2	1.3 versicolor
## 98	6.2	2.9	4.3	1.3 versicolor
## 99	5.1	2.5	3.0	1.1 versicolor
## 100	5.7	2.8	4.1	1.3 versicolor
## 101	6.3	3.3	6.0	2.5 virginica
## 102	5.8	2.7	5.1	1.9 virginica
## 103	7.1	3.0	5.9	2.1 virginica
## 104	6.3	2.9	5.6	1.8 virginica
## 105	6.5	3.0	5.8	2.2 virginica
## 106	7.6	3.0	6.6	2.1 virginica
## 107	4.9	2.5	4.5	1.7 virginica
## 108	7.3	2.9	6.3	1.8 virginica
## 109	6.7	2.5	5.8	1.8 virginica
## 110	7.2	3.6	6.1	2.5 virginica
## 111	6.5	3.2	5.1	2.0 virginica
## 112	6.4	2.7	5.3	1.9 virginica
## 113	6.8	3.0	5.5	2.1 virginica
## 114	5.7	2.5	5.0	2.0 virginica
## 115	5.8	2.8	5.1	2.4 virginica
## 116	6.4	3.2	5.3	2.3 virginica
## 117	6.5	3.0	5.5	1.8 virginica
## 118	7.7	3.8	6.7	2.2 virginica
## 119	7.7	2.6	6.9	2.3 virginica
## 120	6.0	2.2	5.0	1.5 virginica
## 121	6.9	3.2	5.7	2.3 virginica
## 122	5.6	2.8	4.9	2.0 virginica
## 123	7.7	2.8	6.7	2.0 virginica
## 124	6.3	2.7	4.9	1.8 virginica
## 125	6.7	3.3	5.7	2.1 virginica
## 126	7.2	3.2	6.0	1.8 virginica
## 127	6.2	2.8	4.8	1.8 virginica
## 128	6.1	3.0	4.9	1.8 virginica
## 129	6.4	2.8	5.6	2.1 virginica
## 130	7.2	3.0	5.8	1.6 virginica
## 131	7.4	2.8	6.1	1.9 virginica
## 132	7.9	3.8	6.4	2.0 virginica
## 133	6.4	2.8	5.6	2.2 virginica
## 134	6.3	2.8	5.1	1.5 virginica
## 135	6.1	2.6	5.6	1.4 virginica
## 136	7.7	3.0	6.1	2.3 virginica
## 137	6.3	3.4	5.6	2.4 virginica

```
## 138      6.4      3.1      5.5      1.8 virginica
## 139      6.0      3.0      4.8      1.8 virginica
## 140      6.9      3.1      5.4      2.1 virginica
## 141      6.7      3.1      5.6      2.4 virginica
## 142      6.9      3.1      5.1      2.3 virginica
## 143      5.8      2.7      5.1      1.9 virginica
## 144      6.8      3.2      5.9      2.3 virginica
## 145      6.7      3.3      5.7      2.5 virginica
## 146      6.7      3.0      5.2      2.3 virginica
## 147      6.3      2.5      5.0      1.9 virginica
## 148      6.5      3.0      5.2      2.0 virginica
## 149      6.2      3.4      5.4      2.3 virginica
## 150      5.9      3.0      5.1      1.8 virginica
```

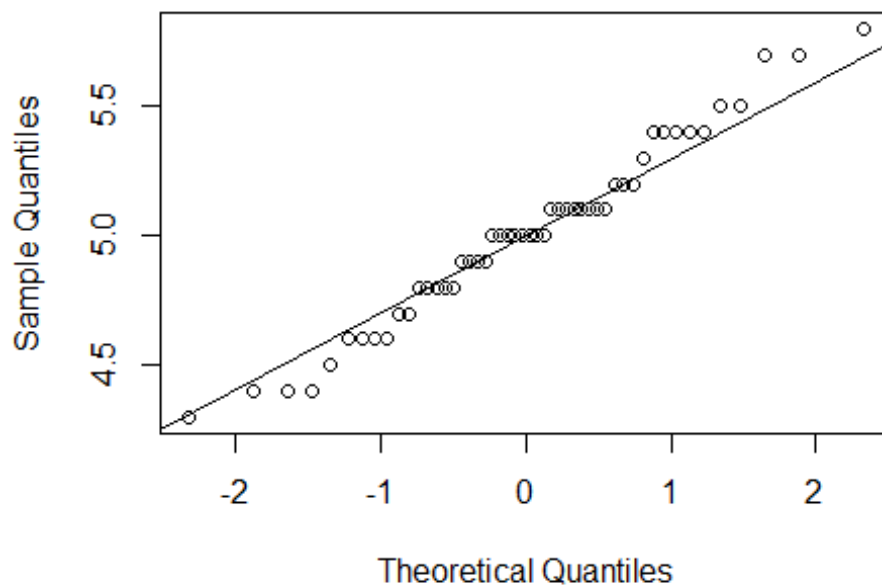
```
sort1 <- iris$Sepal.Length[iris$Species == 'setosa']

sort2 <- iris$Sepal.Length[iris$Species == 'versicolor']

sort3 <- iris$Sepal.Length[iris$Species == 'virginica']

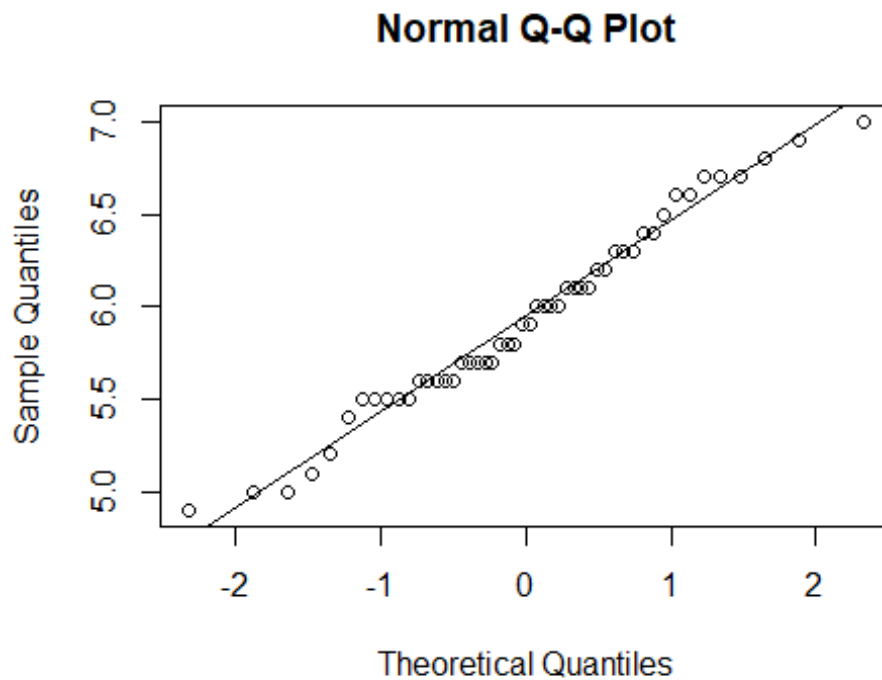
#checking whether the data is normal distributed
qqnorm(sort1)
qqline(sort1)
```

Normal Q-Q Plot

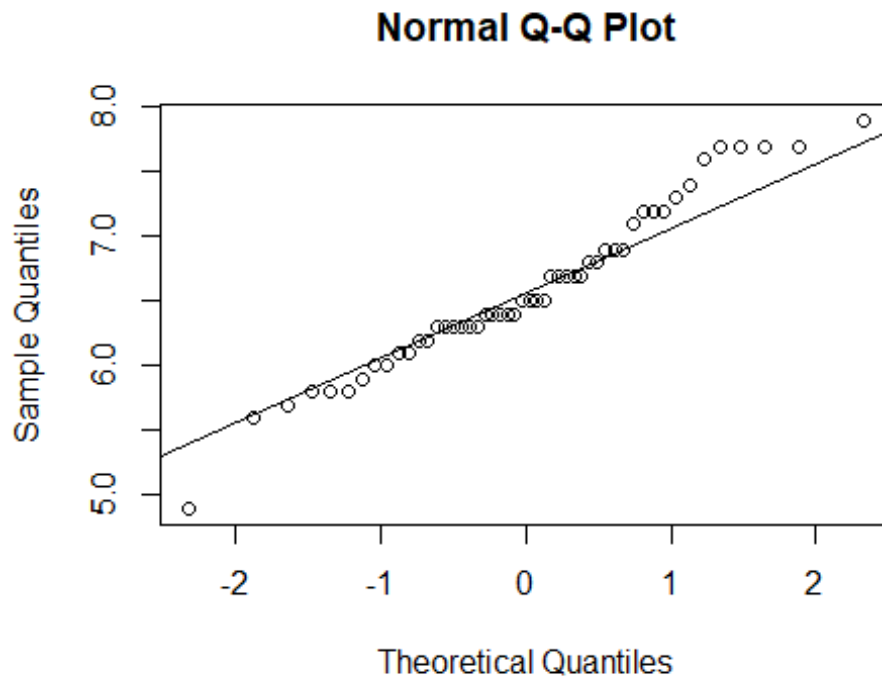


```
shapiro.test(sort1)
```

```
##  
##  Shapiro-Wilk normality test  
##  
## data:  sort1  
## W = 0.9777, p-value = 0.4595  
  
qqnorm(sort2)  
qqline(sort2)
```



```
shapiro.test(sort2)  
  
##  
##  Shapiro-Wilk normality test  
##  
## data:  sort2  
## W = 0.97784, p-value = 0.4647  
  
qqnorm(sort3)  
qqline(sort3)
```

```
shapiro.test(sort3)

##
##  Shapiro-Wilk normality test
##
## data:  sort3
## W = 0.97118, p-value = 0.2583

#all three are normally distributed

# формула на модела (имаме два отклика)
(cbind(iris$Sepal.Length, iris$Sepal.Width) ~ Species) %>%
  # изпълнение на апона с много у променливи
  manova(data = iris) %>%
  # Обобщение
  summary()

##              Df  Pillai approx F num Df den Df    Pr(>F)
## Species       2 0.94531   65.878      4    294 < 2.2e-16 ***
## Residuals 147
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

# Извод: Различните сортове играят роля за размера на чашелистчетата
# някоя от групите има значително различно средно от останалите
```

От изпит 2017

```

#1.
#number of people younger than 20 yrs
length(Aids2$age[Aids2$age < 20])

## [1] 39

#sex of the patients with earliest diagnosis
Aids2$sex[head(order(Aids2$diag), 5)]

## [1] M M M M M
## Levels: F M

#men who got aids from blood
men_blood <- sum(Aids2$sex[Aids2$T.categ == 'blood'] == 'M')

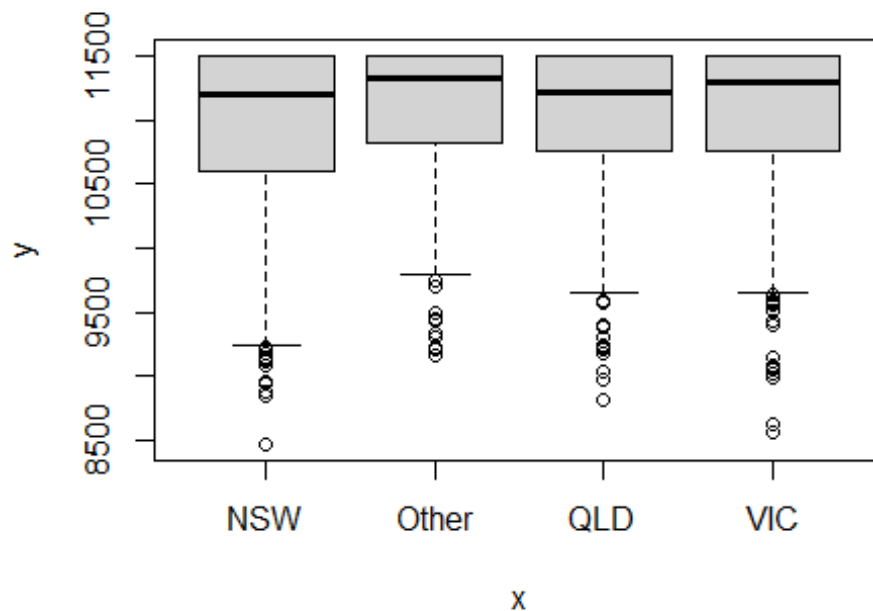
all_men <- sum(Aids2$sex == 'M')

men_blood / all_men

## [1] 0.02069717

#графика за щатът на пациента и смъртността
plot(Aids2$state, Aids2$death, na.rm = T)

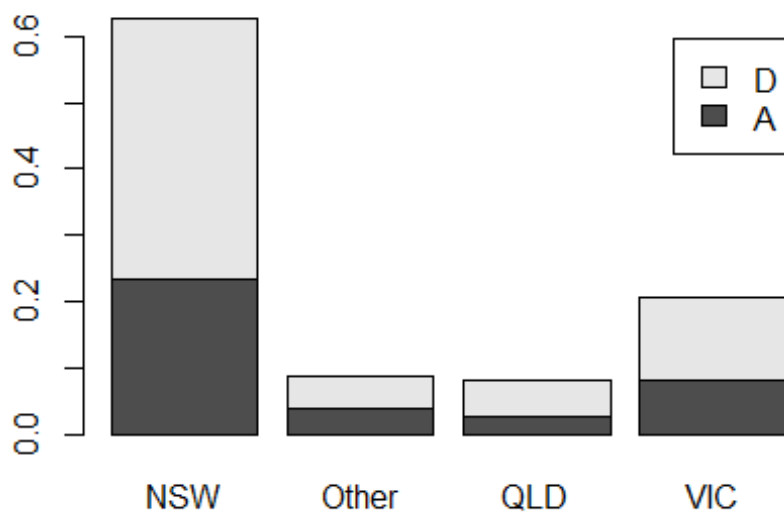
```



```

table(Aids2$status, Aids2$state) %>% prop.table() %>% barplot(legend.text =
T)

```



```
?Aids2
## starting httpd help server ... done

#2.
#total women and women that died
women <- sum(Aids2$sex == 'F')
dead_women <- sum(Aids2$status[Aids2$sex == 'F'] == 'D')

men <- sum(Aids2$sex == 'M')
dead_men <- sum(Aids2$status[Aids2$sex == 'M'] == 'D')

#X0 - жените умират по-малко X1 - умират повече
prop.test(c(dead_women, dead_men), c(women, men), alternative = 'greater')

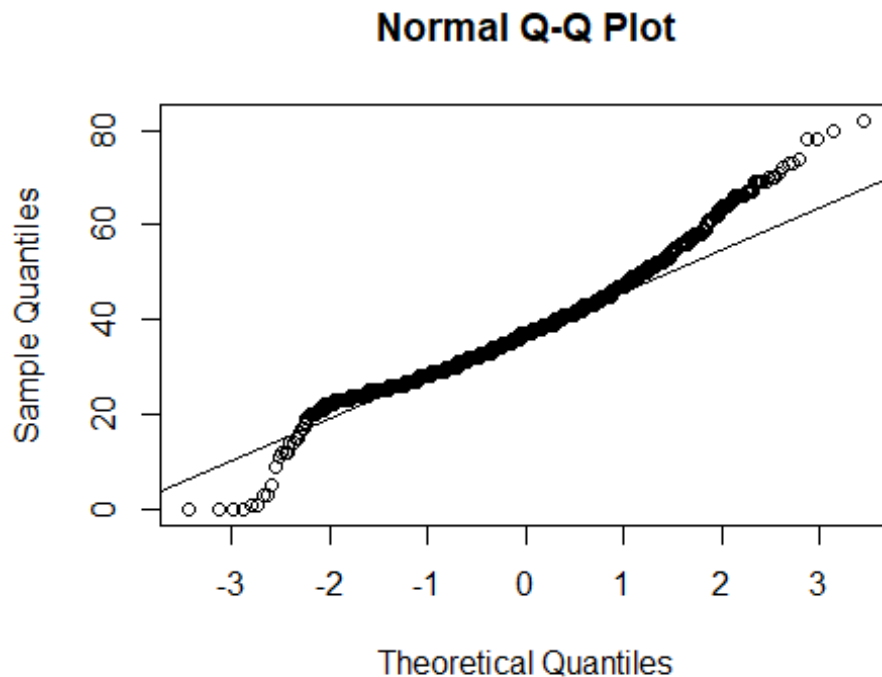
##
## 2-sample test for equality of proportions with continuity correction
##
## data:  c(dead_women, dead_men) out of c(women, men)
## X-squared = 0.13041, df = 1, p-value = 0.641
## alternative hypothesis: greater
## 95 percent confidence interval:
## -0.1173963 1.0000000
## sample estimates:
##  prop 1    prop 2
## 0.5955056 0.6201888
```

```
#можем да приемем нулевата хипотеза
```

```
number_dead <- Aids2$age[Aids2$status == 'D']
```

```
qqnorm(number_dead)
```

```
qqline(number_dead)
```



```
shapiro.test(number_dead)
```

```
##
```

```
## Shapiro-Wilk normality test
```

```
##
```

```
## data: number_dead
```

```
## W = 0.96911, p-value < 2.2e-16
```

```
#not normally distributed
```

```
#X0 - средната възраст е 38 X1 - не е
```

```
#приемаме h1 хипотеза
```

```
wilcox.test(number_dead, mu = 38, alternative = 'two.sided')
```

```
##
```

```
## Wilcoxon signed rank test with continuity correction
```

```
##
```

```
## data: number_dead
```

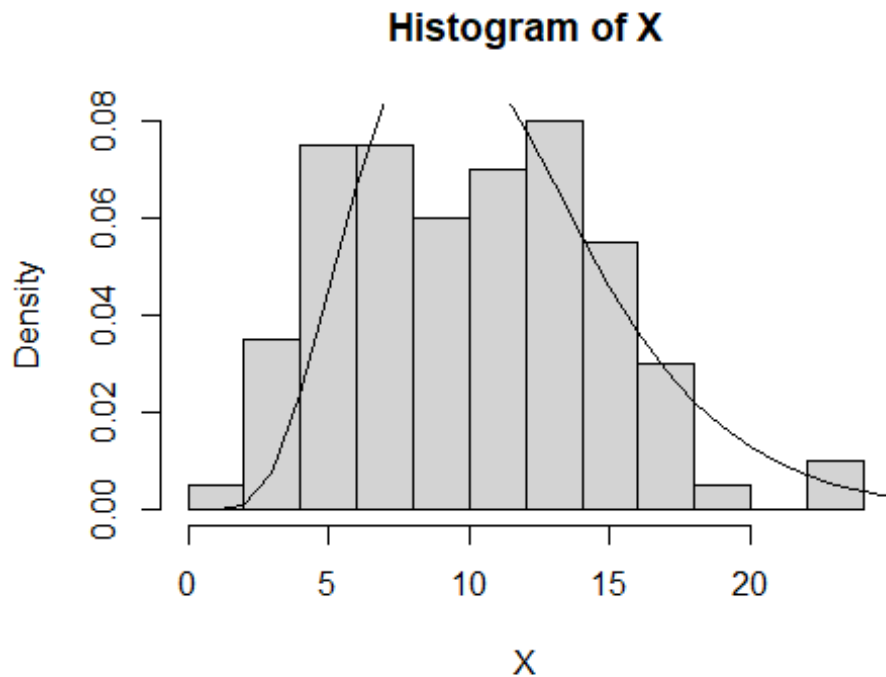
```
## V = 649726, p-value = 0.00164
```

```
## alternative hypothesis: true location is not equal to 38
```

```
#4.
X <- rchisq(100, df = 10)

hist(X, probability = T)

lines(dchisq(0:30, df = 10))
```



```
#5.
cat <- cats[cats$Sex == 'M', ]

#create the model
s <- lm(Hwt ~ Bwt, data = cat) %>% summary()
#based on the model we get that the heart and body weight are not independant
#демек са зависими и сърцето се повлиява от теглото на котката

t <- (s$coefficients[2, 1] - 5) / s$coefficients[2, 2]
#t is negative => we calculate p-value like this:
#df is equal to: numOfObservations - numOfEvaluatedParameters - 1
# so: 97 - 2 - 1 = 94
#Вярно ли е, че при котки по тежки с 1 кг сърцето е по тежко с 5 гр - H0
pval = 2 * pt(t, df = 94)
#тук имаме p-value < 0.05 което значи че отхвърляме нулевата хипотеза, т.е
горното не е вярно

# pval = 2 * pt(t, df = 94, lower.tail = F) if t > 0 only lower tail
```

```
#we test if the distribution is normal so we can do t.test to get the conf.interval
```

```
shapiro.test(cat$Hwt[cat$Bwt == 2.6])
```

```
##
```

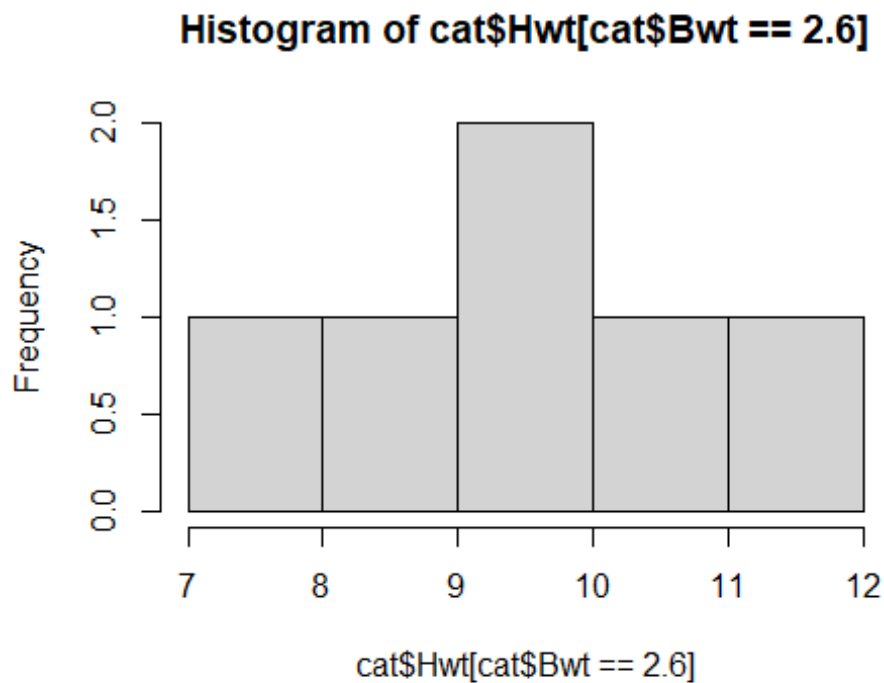
```
## Shapiro-Wilk normality test
```

```
##
```

```
## data: cat$Hwt[cat$Bwt == 2.6]
```

```
## W = 0.96653, p-value = 0.8683
```

```
hist(cat$Hwt[cat$Bwt == 2.6])
```



```
#it is norm dist so we use t.test else we use wilcox.test()
```

```
t.test(cat$Hwt[cat$Bwt == 2.6], conf.level = 0.95)
```

```
##
```

```
## One Sample t-test
```

```
##
```

```
## data: cat$Hwt[cat$Bwt == 2.6]
```

```
## t = 16.654, df = 5, p-value = 1.426e-05
```

```
## alternative hypothesis: true mean is not equal to 0
```

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

```
## 8.005474 10.927859
```

```
## sample estimates:
```

```
## mean of x
```

```
## 9.466667
```

От примерен тест 2017

```

#1.
qnorm(p = 0.05)

## [1] -1.644854

#2.
nrow(state.x77)

## [1] 50

#подреждаме щатите по ниво на необразованост
dumb_states <- head(order(state.x77[,3], decreasing = T), 5)
#взимаме ги според индексите на първите 5 щата
state.x77[dumb_states, 3]

##      Louisiana      Mississippi South Carolina      New Mexico      Texas
##           2.8           2.4           2.3           2.2           2.2

#states with life expectancy over 70
old_states <- state.x77[1:50, 4] > 70
length(state.x77[old_states, 4])

## [1] 41

#щат с най-голяма гъстота на населението
pop <- state.x77[1:50,1]

land <- state.x77[1:50, 8]

density <- pop / land

density[head(order(density, decreasing = T), 1)]

## New Jersey
## 0.9750033

#общото население на петте най-големи щати
biggest_states <- head(order(state.x77[1:50, 8], decreasing = T), 5)
sum(state.x77[biggest_states, 1])

## [1] 35690

#3.но - има по-малко подобрили се жени x1- има повече подобрили се мъже
women <- 200
men <- 100

not_accepted_women <- women * 38 / 100
not_accepted_men <- men * 50 / 100

#приемаме хипотезата че е по-ефективно при жените отколкото при мъжете
prop.test(c(not_accepted_women, not_accepted_men), c(women, men), alternative
= 'greater' )

```

```
##
## 2-sample test for equality of proportions with continuity correction
##
## data:  c(not_accepted_women, not_accepted_men) out of c(women, men)
## X-squared = 3.4637, df = 1, p-value = 0.9686
## alternative hypothesis: greater
## 95 percent confidence interval:
## -0.2272546  1.0000000
## sample estimates:
## prop 1 prop 2
##  0.38  0.50
```

#4.

```
data <- data.frame(
  anscombe$x3,
  anscombe$x4)

l <- lm(data$anscombe.x3 ~ data$anscombe.x4, data = data)

plot(data$anscombe.x3, data$anscombe.x4)
abline(l, col = "red ", lwd = 2)
```

