

Modele regresji i ich zastosowania

Labolatoria 7 i 8

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7 maja 2021

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1 Zadania laboratoryjne

1.1 Zadanie 1

Generujemy $n = 100$ obserwacji Y_1, \dots, Y_n postaci

$$Y_i = \beta_1 + \beta_2 e^{-\beta_3 x} + \epsilon_i \quad (1)$$

gdzie $\epsilon_1, \dots, \epsilon_n$ i.i.d. $N(0, \sigma^2)$

```
n = 100
set.seed(100)
funkcja1 <- function(i) {
  x_i <- 10*i/n
  return(x_i)
}
```

```
x_100<-seq(1,100)
```

```
x_i<-funkcja1(x_100)
```

```
beta1 = 80
```

```

beta2 = 100
beta3 = 0.005
sigma2 = 0.5

epsilon_i<-rnorm(100,0,sigma2)
exp(-beta3*0.1)

## [1] 0.9995001

y_i<-beta1+beta2*exp(-beta3*x_i)+epsilon_i

head(cbind(x_i,y_i))

##      x_i      y_i
## [1,] 0.1 179.6989
## [2,] 0.2 179.9658
## [3,] 0.3 179.8107
## [4,] 0.4 180.2436
## [5,] 0.5 179.8088
## [6,] 0.6 179.8598

```

1.2 Zadanie 2

Znajdujemy postać funkcji $g(x, \beta)$ oraz kolejno elementy gradientu G , które są pochodnymi cząstkowymi:

$$g(x, \beta) = \beta_1 + \beta_2 e^{-\beta_3 x} \quad (2)$$

$$\frac{\delta g(x, \beta_1)}{\delta \beta_1} = 1 \quad (3)$$

$$\frac{\delta g(x, \beta_2)}{\delta \beta_2} = e^{-\beta_3 x} \quad (4)$$

$$\frac{\delta g(x, \beta_3)}{\delta \beta_3} = -x \beta_2 e^{-\beta_3 x} \quad (5)$$

1.3 Zadanie 3 i 4

Implementujemy algorytm Gaussa- Newtona zadając parametry wstępne 79,101,0.004. Algorytm przerywa estymacje kolejnych parametrów po spełnieniu kryterium stopu. Tworzymy tabele ze wszystkimi wykonanymi krokami, gdzie liczona jest również sigma oraz wartość log-likelihood.

```

install.packages("MASS")

## Error in contrib.url(repos, "source"): trying to use CRAN without setting a mirror

library(MASS)
matrix0<-matrix(rep(0,5000),ncol=5,nrow=1000)
beta0 <- c(79.0000000,101.0000000,0.0040000)

```

```

for(i in 1:1000) {
  e = y_i - beta0[1] - beta0[2]*exp(-beta0[3]*x_i)
  sigma<-(1/n)*(sum(e^2))
  likelihood<-(-n/2)*log(2*pi)-n/2*log(sigma)-n/2
  matrix0[i,]<-c(beta0,sigma,likelihood)

  gradient = cbind(rep(1,n), exp(-beta0[3]*x_i), -beta0[2]*x_i*exp(-beta0[3]*x_i))
  beta1<-beta0 + ginv(t(gradient)%*%gradient)%*%t(gradient)%*%e

  if(norm(beta0-beta1, type="2")<=0.0001){
    matrix0[i+1,]<-c(beta1,sigma,likelihood)
    break
  }
  else{
    beta0<-beta1
  }
}
head(matrix0)

##           [,1]      [,2]      [,3]      [,4]      [,5]
## [1,] 79.00000 101.0000 0.004000000 0.5676750 -113.58354
## [2,] 79.03417 101.0342 0.005079991 0.2563989 -73.84280
## [3,] 79.03462 101.0346 0.005085324 0.2563936 -73.84178
## [4,] 79.03462 101.0346 0.005085323 0.2563936 -73.84178
## [5,] 0.00000 0.0000 0.000000000 0.0000000 0.00000
## [6,] 0.00000 0.0000 0.000000000 0.0000000 0.00000

matrix0<-matrix0[1:4,1:5]
matrix0

##           [,1]      [,2]      [,3]      [,4]      [,5]
## [1,] 79.00000 101.0000 0.004000000 0.5676750 -113.58354
## [2,] 79.03417 101.0342 0.005079991 0.2563989 -73.84280
## [3,] 79.03462 101.0346 0.005085324 0.2563936 -73.84178
## [4,] 79.03462 101.0346 0.005085323 0.2563936 -73.84178

colnames(matrix0)<-c("beta_1","beta_2","beta_3","sigma","log-likelihood")
row.names(matrix0)<-c(0,1,2,3)
matrix0

##      beta_1  beta_2      beta_3      sigma log-likelihood
## 0 79.00000 101.0000 0.004000000 0.5676750      -113.58354
## 1 79.03417 101.0342 0.005079991 0.2563989      -73.84280
## 2 79.03462 101.0346 0.005085324 0.2563936      -73.84178
## 3 79.03462 101.0346 0.005085323 0.2563936      -73.84178

```

- Zostały wykonane 3 iteracje.
- Widzimy bardzo niskie różnice między kolejnymi estymacjami
- Krok 3 nie ma już sensu

1.4 Zadanie 5

Porównanie parametrów, różnice

```
beta000 <- c(79.0000000,101.0000000,0.0040000)
matrix0[3,1:3]-beta000
```

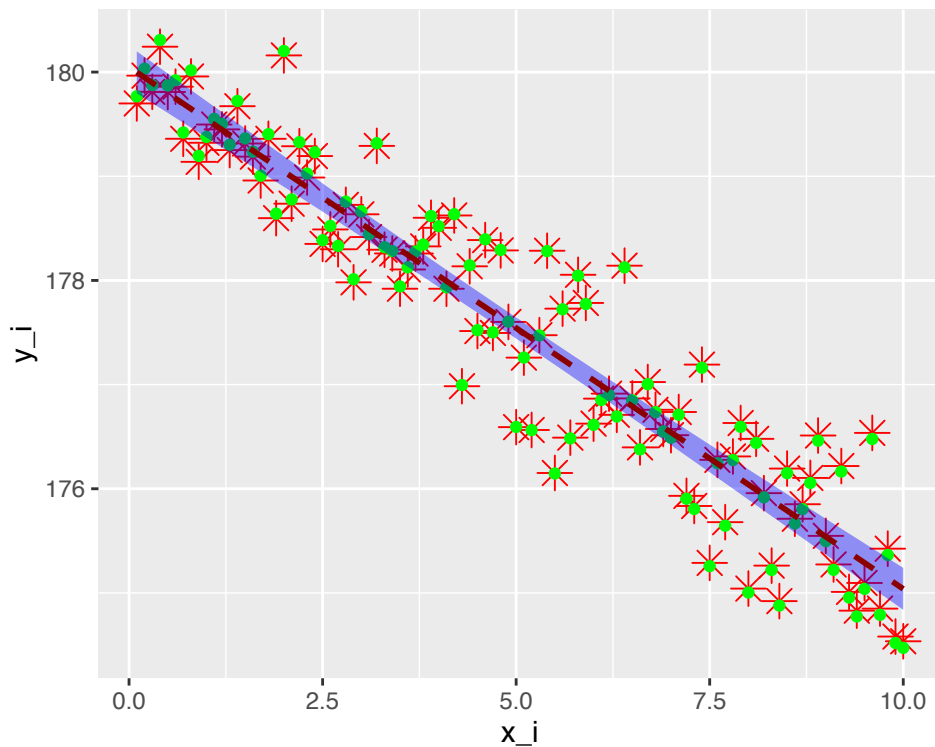
```
##      beta_1      beta_2      beta_3
## 0.034617830 0.034622669 0.001085324
```

1.5 Zadanie 6

```
beta_00<-matrix0[4,1:3]
y_i_2<-beta_00[1]+beta_00[2]*exp(-beta_00[3]*x_i)+epsilon_i
y<-c(y_i,y_i_2)
```

```
library(ggplot2)
ggplot(data.frame(x_i,y_i), aes(x = x_i, y = y_i)) +
  geom_point(shape=8, color="red", lwd =4)+
  geom_point(data = data.frame(x_i,y_i_2), aes(x = x_i, y = y_i_2),col = "green")+
  geom_smooth(method=lm, linetype="dashed",
              color="darkred", fill="blue")
```

```
## 'geom_smooth()' using formula 'y ~ x'
```



- Widzimy, że punkty niemal się całkowicie pokrywają na wykresie

1.6 Zadanie 7

Macierz kowariancji

```
cov_matrix<-data.frame(matrix0[1:3,1],matrix0[1:3,2],matrix0[1:3,3])
cov(cov_matrix)
```

```
##                matrix0.1.3..1. matrix0.1.3..2. matrix0.1.3..3.
## matrix0.1.3..1.    3.943752e-04    3.944301e-04    1.241305e-05
## matrix0.1.3..2.    3.944301e-04    3.944851e-04    1.241478e-05
## matrix0.1.3..3.    1.241305e-05    1.241478e-05    3.907229e-07
```

Macierz kowariancji jest bliska macierzy zerowej.

1.7 Zadanie 8

Parametry nieznacznie różniące się od β początkowej

```
matrix0<-matrix(rep(0,5000),ncol=5,nrow=1000)
beta0 <- c(82,102,0.006)
for(i in 1:1000) {
  e = y_i - beta0[1] - beta0[2]*exp(-beta0[3]*x_i)
  sigma<-(1/n)*(sum(e^2))
  likelihood<-(-n/2)*log(2*pi)-(n/2)*log(sigma)-n/2
  matrix0[i,]<-c(beta0,sigma,likelihood)
```

```

gradient = cbind(rep(1,n), exp(-beta0[3]*x_i), -beta0[2]*x_i*exp(-beta0[3]*x_i))
beta1<-beta0 + ginv(t(gradient)%*%gradient)%*%t(gradient)%*%e

if(norm(beta0-beta1, type="2")<=0.0001){
  matrix0[i+1,]<-c(beta1,sigma,likelihood)
  break
}
else{
  beta0<-beta1
}
}
matrix0<-matrix0[1:4,1:5]
colnames(matrix0)<-c("beta_1","beta_2","beta_3","sigma","log-likelihood")
row.names(matrix0)<-c(0,1,2,3)
matrix0

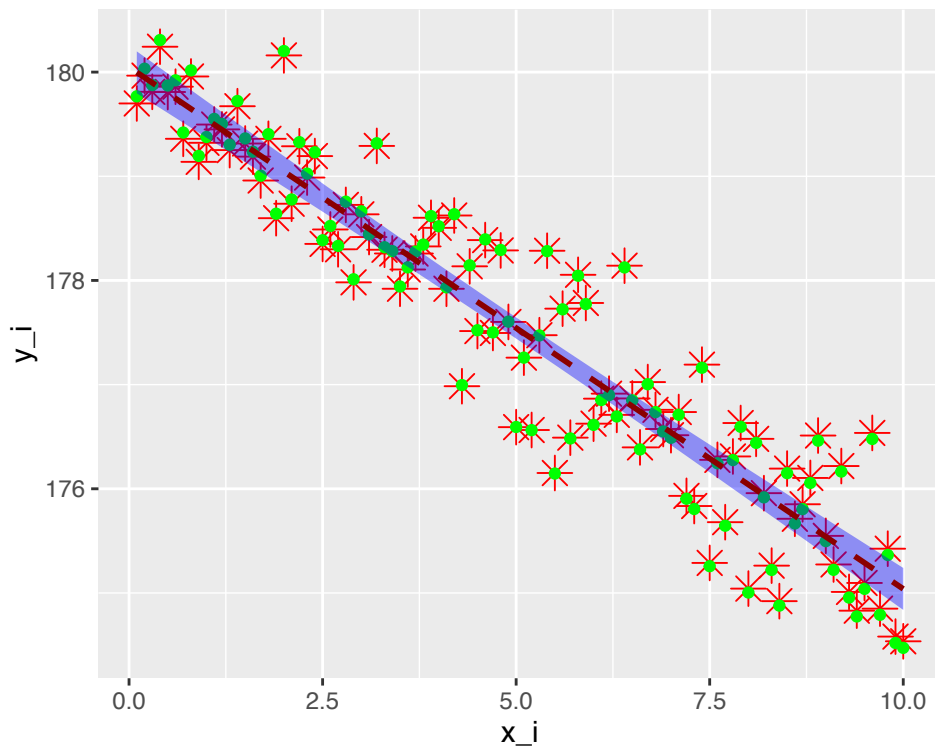
##      beta_1  beta_2      beta_3      sigma log-likelihood
## 0 82.00000 102.0000 0.006000000 12.2473392      -267.15929
## 1 80.03468 100.0341 0.005150263  0.2564581      -73.85435
## 2 80.03503 100.0344 0.005137497  0.2563978      -73.84260
## 3 80.03503 100.0344 0.005137500  0.2563978      -73.84260

beta_00<-matrix0[4,1:3]
y_i_2<-beta_00[1]+beta_00[2]*exp(-beta_00[3]*x_i)+epsilon_i
y<-c(y_i,y_i_2)

ggplot(data.frame(x_i,y_i), aes(x = x_i, y = y_i)) +
  geom_point(shape=8, color="red", lwd =4)+
  geom_point(data = data.frame(x_i,y_i_2), aes(x = x_i, y = y_i_2),col = "green")+
  geom_smooth(method=lm, linetype="dashed",
              color="darkred", fill="blue")

## 'geom_smooth()' using formula 'y ~ x'

```



```

matrix0<-matrix(rep(0,5000),ncol=5,nrow=1000)
beta0 <- c(43,180,0.01)
for(i in 1:1000) {
  e = y_i - beta0[1] - beta0[2]*exp(-beta0[3]*x_i)
  sigma<-(1/n)*(sum(e^2))
  likelihood<-(-n/2)*log(2*pi)-(n/2)*log(sigma)-n/2
  matrix0[i,]<-c(beta0,sigma,likelihood)

  gradient = cbind(rep(1,n), exp(-beta0[3]*x_i), -beta0[2]*x_i*exp(-beta0[3]*x_i))
  beta1<-beta0 + ginv(t(gradient)%*%gradient)%*%t(gradient)%*%e

  if(norm(beta0-beta1, type="2")<=0.0001){
    matrix0[i+1,]<-c(beta1,sigma,likelihood)
    break
  }
  else{
    beta0<-beta1
  }
}
matrix0<-matrix0[1:5,1:5]
colnames(matrix0)<-c("beta_1","beta_2","beta_3","sigma","log-likelihood")
row.names(matrix0)<-c(0,1,2,3,4)
matrix0

##      beta_1  beta_2      beta_3      sigma log-likelihood

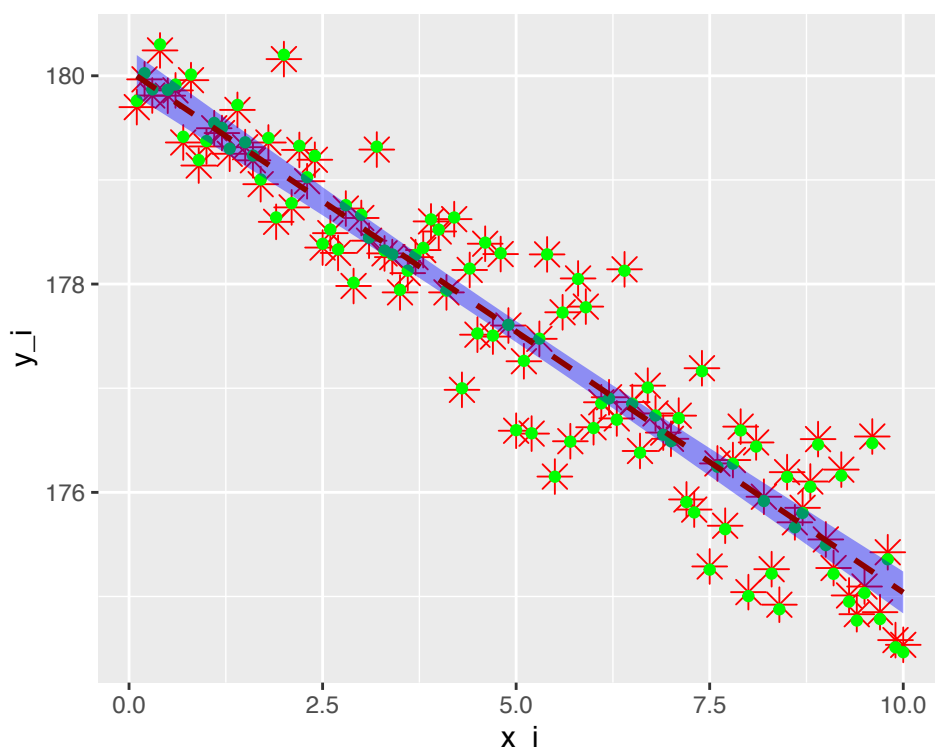
```

```
## 0 43.00000 180.0000 0.010000000 1358.5466047 -502.60239
## 1 21.50864 158.4902 0.003812235 0.6096497 -117.15031
## 2 21.53964 158.5212 0.003209045 0.2562568 -73.81509
## 3 21.53991 158.5214 0.003211053 0.2562549 -73.81473
## 4 21.53991 158.5214 0.003211053 0.2562549 -73.81473

beta_00<-matrix0[4,1:3]
y_i_2<-beta_00[1]+beta_00[2]*exp(-beta_00[3]*x_i)+epsilon_i
y<-c(y_i,y_i_2)

ggplot(data.frame(x_i,y_i), aes(x = x_i, y = y_i)) +
  geom_point(shape=8, color="red", lwd =4)+
  geom_point(data = data.frame(x_i,y_i_2), aes(x = x_i, y = y_i_2), col = "green")+
  geom_smooth(method=lm, linetype="dashed",
              color="darkred", fill="blue")

## 'geom_smooth()' using formula 'y ~ x'
```



- Widzimy, że zarówno dla małych jak i dużych rozbieżności parametrów wstępnych za- uważać możemy szybkie dopasowanie estymatorów (w 3 i 4 krokach) oraz bardzo bliskie dopasowanie na wykresach.