

# Computational Statistics. Chapter 3: EM algorithm. Solution of exercises

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## Exercise 1

### Question 1a

We first give values to the model parameters:

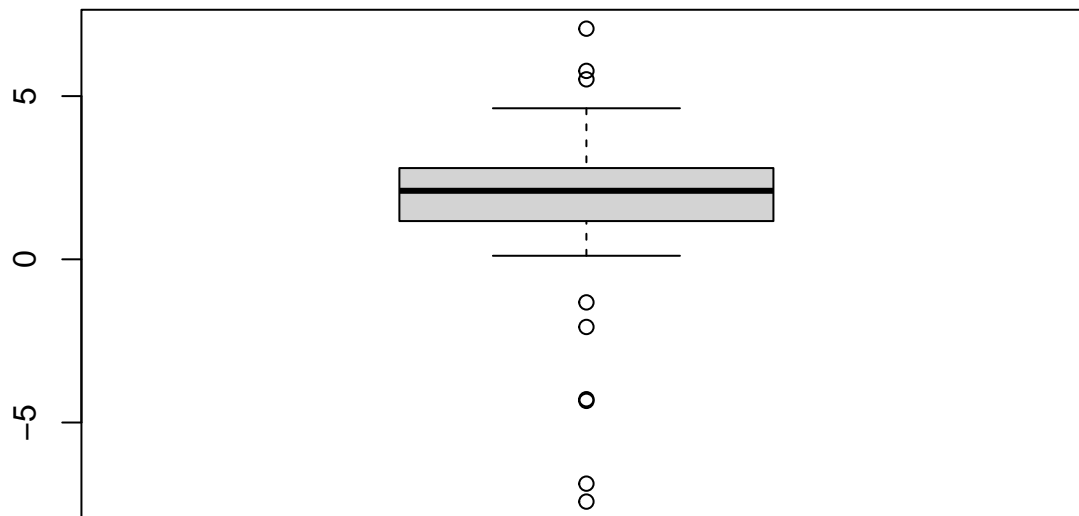
```
pi <- 0.90
mu <- 2
sig <- 1
a <- 10
c <- 1/(2*a)
n <- 100
```

We then generate the data:

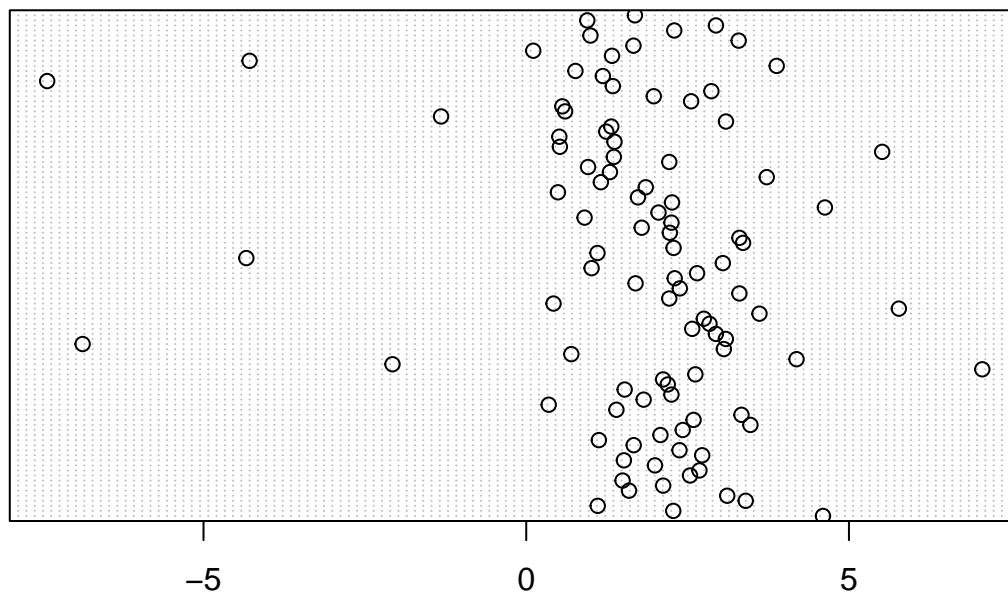
```
y<-vector("numeric",n)
z<-vector("numeric",n)
for(i in 1:n){
  z[i] <- sample(c(1,0),size=1,prob=c(pi,1-pi))
  if(z[i]==1)
    y[i] <- rnorm(1,mean=mu,sd=sig)
  else y[i] <- runif(1,min=-a,max=a)
}
```

Finally, we generate box and dot plots the data:

```
boxplot(y)
```



```
dotchart(y)
```



## Question 1b

We first write a function that computes the observed-data log-likelihood:

```
loglik<- function(theta,y){
  phi <- sapply(y,dnorm,mean=theta[1],sd=theta[2])
  logL <- sum(log(theta[3]*phi+(1-theta[3])*c))
  return(logL)
}
```

We then write the EM algorithm for this problem. The inputs are the data  $y$ , the initial parameter value  $\theta_0$ , the constant  $a$ , the threshold  $\epsilon$  used in the stopping criterion, and a flag `disp` that controls the display of the intermediate results. The outputs are the maximum observed-data log-likelihood, the corresponding MLE of  $\theta$ , and the vector  $z$  of estimated probabilities.

```

em_outlier <- function(y,theta0,a,epsi,disp=TRUE){
  go_on<-TRUE
  logL0 <- loglik(theta0,y)
  t<-0
  c<-1/(2*a)
  n<-length(y)
  if(disp) print(c(t,logL0))
  while(go_on){
    t<-t+1
    # E-step
    phi <- sapply(y,dnorm,mean=theta0[1],sd=theta0[2])
    z<- phi*theta0[3]/(phi*theta0[3]+c*(1-theta0[3]))
    # M-step
    S<- sum(z)
    pi<-S/n
    mu<- sum(y*z)/S
    sig<-sqrt(sum(z*(y-mu)^2)/S)
    theta<-c(mu,sig,pi)
    logL<-loglik(theta,y)
    if (logL-logL0 < epsi) go_on <- FALSE
    logL0 <- logL
    theta0<-theta
    if(disp) print(c(t,logL))
  }
  return(list(loglik=logL,theta=theta,z=z))
}

```

## Question 1c

Let us now run the above function with our data. We initialize parameters  $\mu$  and  $\sigma$  with the mean and standard deviation of the data, and we set  $\pi_0 = 0.5$ :

```

mu0<-mean(y) # +rnorm(1,mean=0,sd=0.5)
sig0<-sd(y)
pi0<-0.5
theta0<-c(mu0,sig0,pi0)

```

We then run function `em_outlier`:

```

estim<-em_outlier(y,theta0,a,epsi=1e-6)

```

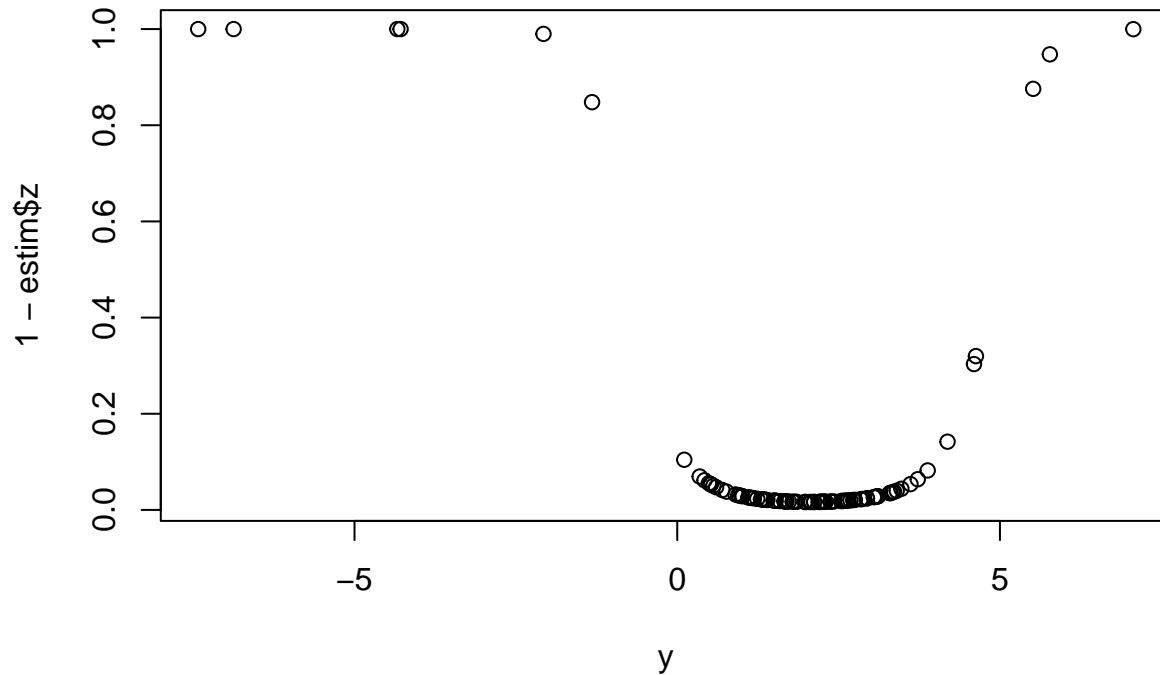
```

## [1] 0.0000 -230.5982
## [1] 1.000 -190.173
## [1] 2.0000 -183.3387
## [1] 3.0000 -182.9011
## [1] 4.0000 -182.8613
## [1] 5.000 -182.854
## [1] 6.0000 -182.8523
## [1] 7.0000 -182.8519
## [1] 8.0000 -182.8518
## [1] 9.0000 -182.8518
## [1] 10.0000 -182.8518
## [1] 11.0000 -182.8518
## [1] 12.0000 -182.8518

```

Finally, we plot the estimated probabilities  $1 - z_i$  against the inputs  $y_i$ :

```
plot(y, 1-estim$z)
```



We can see that the outliers have a high estimated probability of being drawn from the uniform distribution, as expected.

## Exercise 2

### Question 2a

We set the parameters:

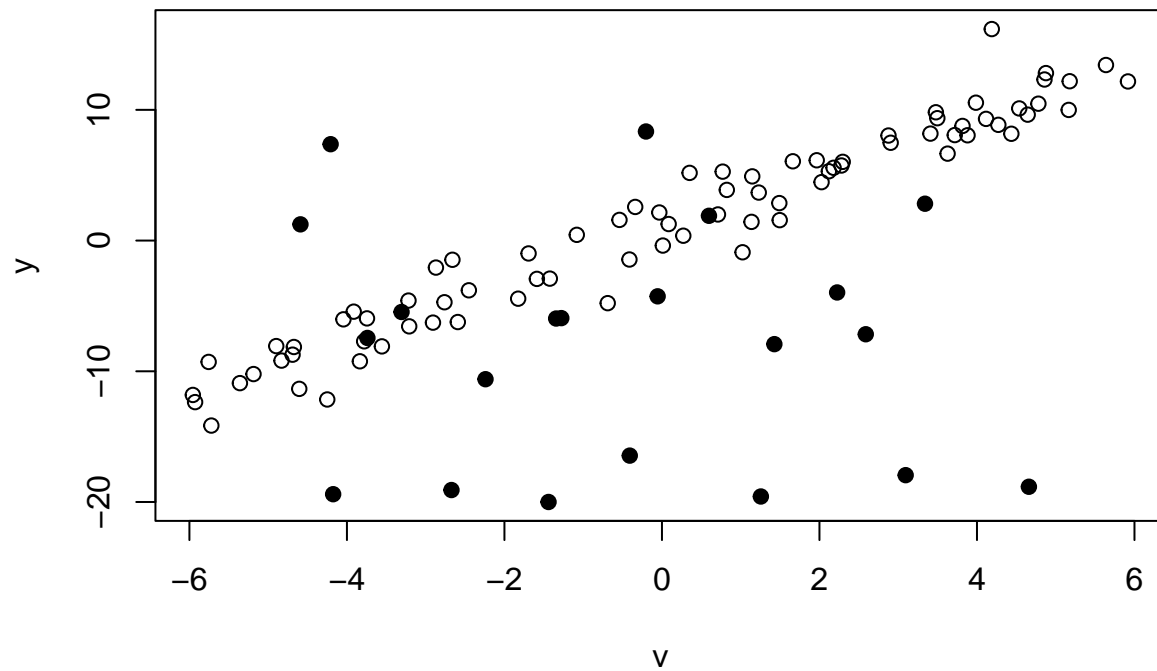
```
pi<-0.8
beta<-c(1,2)
sig<- 2
a<-20
c<-1/(2*a)
n<- 100
```

We then generate the data:

```
y<-vector("numeric",n)
v <- runif(n,min=-6,max=6)
z<-vector("numeric",n)
for(i in 1:n){
  z[i]=sample(c(1,0),size=1,prob=c(pi,1-pi))
  if(z[i]==1)
    y[i]<-rnorm(1,mean=beta[1]+v[i]*beta[2],sd=sig)
  else y[i]<-runif(1,min=-a,max=a)
}
```

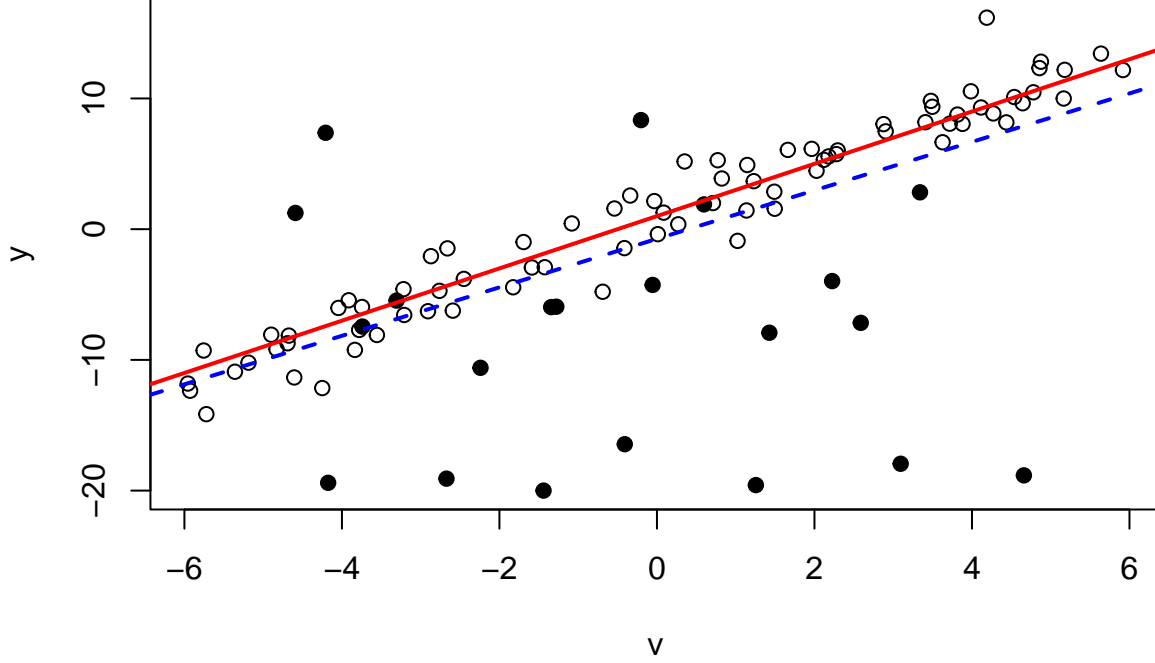
Finally, we plot the data. The data points generated from the uniform distribution (outliers) are highlighted:

```
plot(v,y)
points(v[z==0],y[z==0],pch=16)
```



## Question 2b

```
reg<-lm(y~v)
plot(v,y)
points(v[z==0],y[z==0],pch=16)
abline(reg,lty=2,col="blue",lwd=2) # LS line
abline(1,2,lty=1,col="red",lwd=2) #true regression line
```



### Question 2c

In this problem the observed data is the vector  $\mathbf{y} = (y_1, \dots, y_n)$ . The observed-data likelihood is

$$L(\boldsymbol{\theta}) = \prod_{i=1}^n f(y_i; \boldsymbol{\theta}) = \prod_{i=1}^n [\pi \phi(y_i, v_i^T \beta, \sigma) + (1 - \pi)c].$$

The missing data is the random vector  $\mathbf{Z} = (Z_1, \dots, Z_n)$ , where  $Z_i = 1$  if the observation is not an outlier, and  $Z_i = 0$  otherwise. Clearly,  $Z_i$  has a Bernoulli distribution  $\mathcal{B}(\pi)$ . The complete-data likelihood is

$$L_c(\boldsymbol{\theta}) = \prod_{i=1}^n f(y_i | z_i; \boldsymbol{\theta}) f(z_i; \boldsymbol{\theta}) = \prod_{i=1}^n [\phi(y_i, v_i^T \beta, \sigma)^{z_i} c^{1-z_i} \pi^{z_i} (1 - \pi)^{1-z_i}],$$

and its logarithm is

$$\ell_c(\boldsymbol{\theta}) = \sum_{i=1}^n [z_i \log \phi(y_i, v_i^T \beta, \sigma) + (1 - z_i) \log c + z_i \log \pi + (1 - z_i) \log(1 - \pi)].$$

Function  $Q$  is obtained by replacing the terms  $z_i$  by their conditional expectations given  $\mathbf{Y} = \mathbf{y}$ , for  $\boldsymbol{\theta} = \boldsymbol{\theta}^{(t)}$ . In the E-step, we will compute these conditional expectations as

$$z_i^{(t)} = \mathbb{E}_{\boldsymbol{\theta}^{(t)}}[Z_i | y_i] = \mathbb{P}_{\boldsymbol{\theta}^{(t)}}[Z_i = 1 | y_i] = \frac{\phi(y_i; v_i^T \beta^{(t)}, \sigma^{(t)}) \pi^{(t)}}{\phi(y_i; v_i^T \beta^{(t)}, \sigma^{(t)}) \pi^{(t)} + c(1 - \pi^{(t)})} \quad (1)$$

and

$$\begin{aligned} Q(\boldsymbol{\theta}, \boldsymbol{\theta}^{(t)}) &= \sum_{i=1}^n \left[ z_i^{(t)} \log \phi(y_i, v_i^T \beta, \sigma) + (1 - z_i^{(t)}) \log c + z_i^{(t)} \log \pi + (1 - z_i^{(t)}) \log(1 - \pi) \right] \\ &= \underbrace{\sum_{i=1}^n z_i^{(t)} \log \phi(y_i, v_i^T \beta, \sigma)}_A + \underbrace{\log \pi \sum_{i=1}^n z_i^{(t)} + \log(1 - \pi) \left( n - \sum_{i=1}^n z_i^{(t)} \right)}_B + \underbrace{(1 - z_i^{(t)}) \log c}_C. \end{aligned}$$

In the M-step, we maximize  $Q(\boldsymbol{\theta}, \boldsymbol{\theta}^{(t)})$  with respect to  $\boldsymbol{\theta}$ . In the above equation  $A$  depends only on  $\beta$  and  $\sigma$ ,  $B$  depends on  $\pi$ , and  $C$  is a constant. We can thus maximize  $A$  and  $B$  separately. We have

$$A = -\frac{1}{2} \left( \sum_{i=1}^n z_i^{(t)} \right) \log(2\pi) - \left( \sum_{i=1}^n z_i^{(t)} \right) \log \sigma - \frac{1}{2\sigma^2} \sum_{i=1}^n z_i^{(t)} (y_i - v_i^T \beta)^2.$$

We can see that the vector  $\beta$  maximizing  $A$  (and  $Q$ ) is the solution of a weighted least-squares problem (each individual error term being weighted by  $z_i^{(t)}$ ). We thus have the following update rule for  $\beta$ :

$$\beta^{(t+1)} = \left[ \sum_{i=1}^n z_i^{(t)} v_i v_i^T \right]^{-1} \sum_{i=1}^n z_i^{(t)} v_i y_i.$$

To maximize  $A$  w.r.t.  $\sigma$ , we set  $\beta = \beta^{(t+1)}$  and we compute the derivative:

$$\frac{\partial A}{\partial \sigma} = -\frac{1}{\sigma} \sum_i z_i^{(t)} + \frac{1}{\sigma^3} \sum_i z_i^{(t)} (y_i - v_i^T \beta^{(t+1)})^2.$$

Setting this derivative to zero, we get the update equation for  $\sigma$ :

$$\sigma^{(t+1)} = \sqrt{\frac{\sum_{i=1}^n z_i^{(t)} (y_i - v_i^T \beta^{(t+1)})^2}{\sum_{i=1}^n z_i^{(t)}}}.$$

Finally, to maximize  $B$ , we compute the derivative

$$\frac{\partial B}{\partial \pi} = \frac{\sum_{i=1}^n z_i^{(t)}}{\pi} - \frac{n - \sum_{i=1}^n z_i^{(t)}}{1 - \pi}.$$

Setting this derivative to zero we get the update rule for  $\pi$ :

$$\pi^{(t+1)} = \frac{1}{n} \sum_{i=1}^n z_i^{(t)}.$$

We can now write an EM algorithm for this problem. We first write a function that computes the observed-data log-likelihood:

```
loglik_reg <- function(theta, y, v) {
  n <- length(y)
  phi <- vector(n, mode = "numeric")
  for(i in 1:n)
    phi[i] <- dnorm(y[i], mean = theta[1] + theta[2] * v[i], sd = theta[3])
  logL <- sum(log(theta[4] * phi + (1 - theta[4]) * c))
  return(logL)
}
```

We then write the EM algorithm, which is similar to that of Exercise 1:

```
em_outlier_reg <- function(y, v, theta0, a, epsi) {
  go_on <- TRUE
  logL0 <- loglik_reg(theta0, y, v)
  t <- 0
  c <- 1 / (2 * a)
  n <- length(y)
  phi <- vector(n, mode = "numeric")
  print(c(t, logL0))
```

```

while(go_on){
  t<-t+1
  # E-step
  for(i in 1:n)
    phi[i] <- dnorm(y[i],mean=theta0[1]+theta0[2]*v[i],sd=theta0[3])
  z <- phi*theta0[4]/(phi*theta0[4]+c*(1-theta0[4]))
  # M-step
  S<- sum(z)
  pi<-S/n
  reg<-lm(y ~v,weights=z)
  beta<-reg$coefficients
  sig<-sqrt(sum(z*reg$residuals^2)/S)
  theta<-c(beta,sig,pi)
  logL<-loglik_reg(theta,y,v)
  if (logL-logL0 < epsi) go_on <- FALSE
  logL0 <- logL
  theta0<-theta
  print(c(t,logL))
}
return(list(loglik=logL,theta=theta,z=z))
}

```

## Question 2d

We first initialize the parameters to the OLS estimates and some arbitrary value of  $\pi$ :

```

beta0<-reg$coefficients
sig0<-sd(reg$residuals)
pi0<-0.8
theta0<-c(beta0,sig0,pi0)

```

We then run function `em_outlier_reg`:

```

estim<-em_outlier_reg(y,v,theta0,a,epsi=1e-6)

```

```

## [1] 0.0000 -316.4834
## [1] 1.0000 -288.0812
## [1] 2.0000 -271.7673
## [1] 3.000 -265.862
## [1] 4.0000 -264.6194
## [1] 5.0000 -264.3309
## [1] 6.0000 -264.2446
## [1] 7.0000 -264.2149
## [1] 8.0000 -264.2039
## [1] 9.0000 -264.1997
## [1] 10.000 -264.198
## [1] 11.0000 -264.1973
## [1] 12.0000 -264.1971
## [1] 13.000 -264.197
## [1] 14.0000 -264.1969
## [1] 15.0000 -264.1969
## [1] 16.0000 -264.1969
## [1] 17.0000 -264.1969
## [1] 18.0000 -264.1969

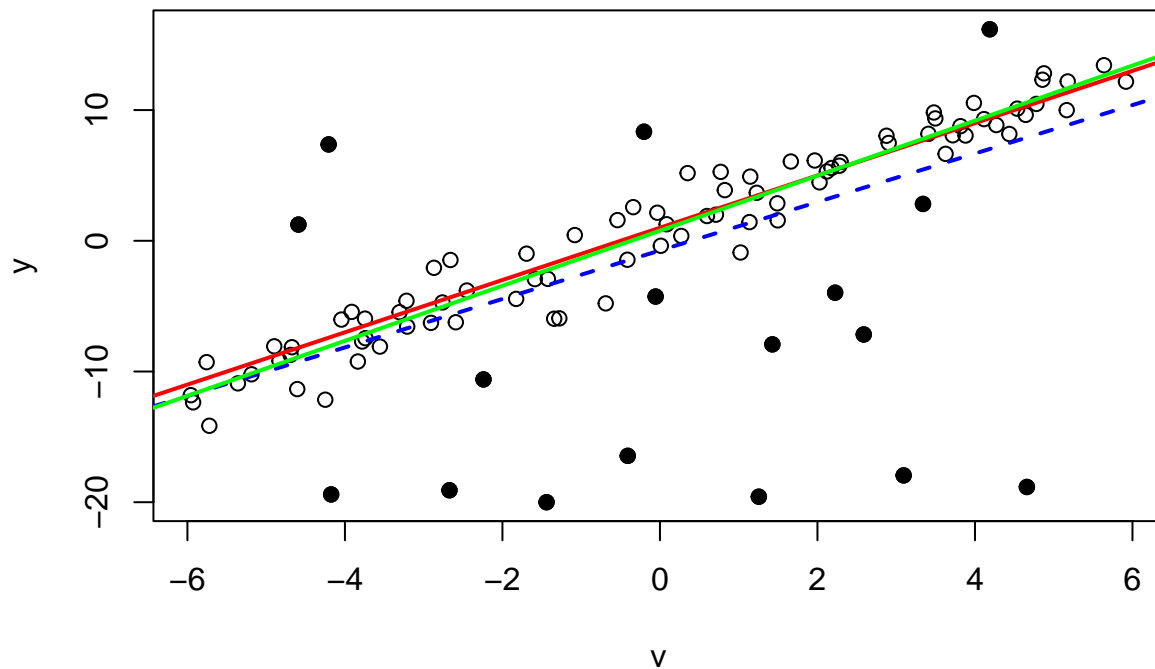
```



```
## [1] 19.0000 -264.1969
```

Finally, we plot the line estimated by EM, together with the true regression line and the OLS line. We also plot the points identified as outliers:

```
plot(v,y)
abline(reg,lty=2,col="blue",lwd=2) # LS line
abline(1,2,lty=1,col="red",lwd=2) # regression line
abline(estim$theta[1],estim$theta[2],lty=1,col='green',lwd=2) # line estimated using EM
points(v[estim$z<0.5],y[estim$z<0.5],pch=19) # points identified as outliers
```



## Exercise 3

### Question 3a

We set the parameters:

```
pi <- 0.8
beta1 <- c(1,2)
sig1 <- 2
beta2 <- c(0,0)
sig2 <- 10
n <- 100
theta_true <- c(beta1,sig1,beta2,sig2,pi)
```

We then generate the data:

```
set.seed(2024022)
y<-vector("numeric",n)
v <- runif(n,min=-6,max=6)
z<-vector("numeric",n)
for(i in 1:n){
  z[i] <- sample(c(1,0),size=1,prob=c(pi,1-pi))
```

```

if(z[i]==1)
  y[i]<- rnorm(1,mean=beta1[1]+v[i]*beta1[2],sd=sig1)
else y[i]<-rnorm(1,mean=beta2[1]+v[i]*beta1[2],sd=sig2)
}

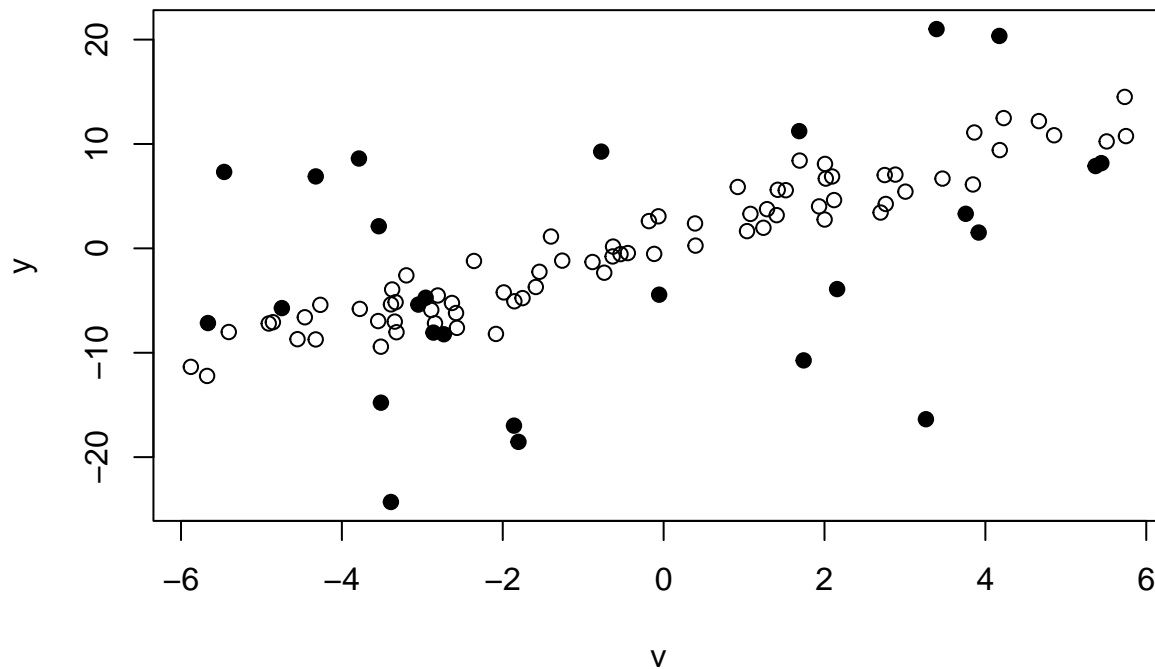
```

Finally, we plot the data. The data points generated from the uniform distribution (outliers) are highlighted:

```

plot(v,y)
points(v[z==0],y[z==0],pch=16)

```

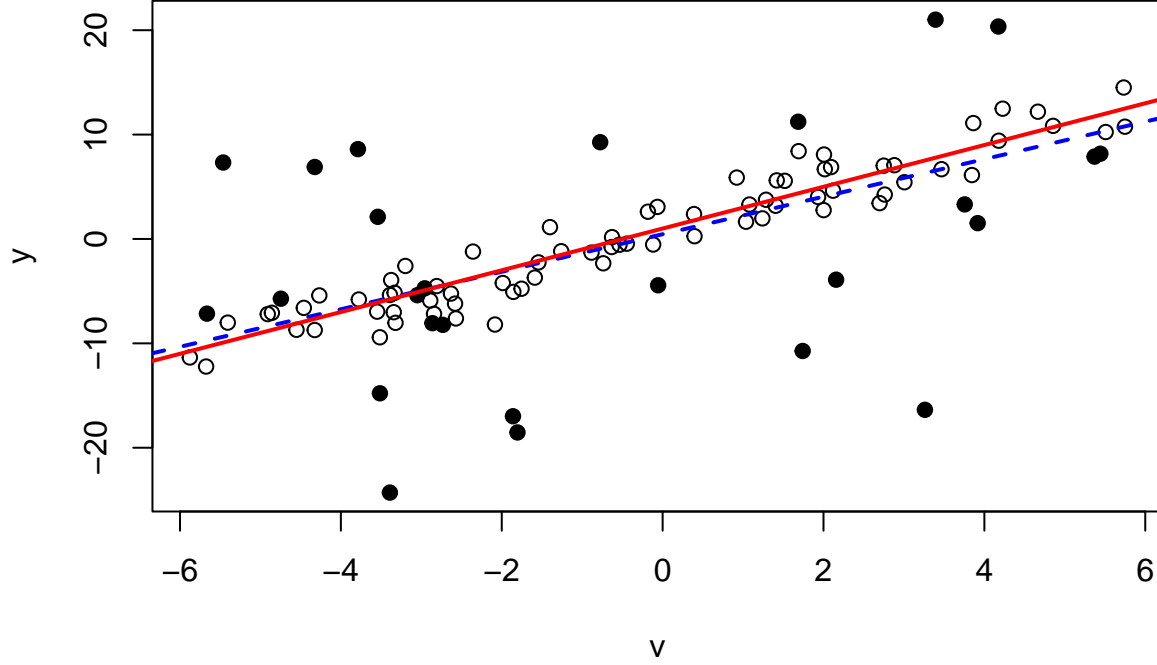


### Question 3b

```

reg<-lm(y~v)
plot(v,y)
points(v[z==0],y[z==0],pch=16)
abline(reg,lty=2,col="blue",lwd=2) # LS line
abline(1,2,lty=1,col="red",lwd=2) #true regression line

```



```
print(sd(reg$residuals))
```

```
## [1] 5.850116
```

We can see that the error standard deviation in the regression model is grossly overestimated because of the outliers.

### Question 3c

The observed-data likelihood is now

$$L(\theta) = \prod_{i=1}^n f(y_i; \theta) = \prod_{i=1}^n [\pi \phi(y_i, v_i^T \beta_1, \sigma_1) + (1 - \pi) \phi(y_i, v_i^T \beta_2, \sigma_2)] .$$

The complete-data likelihood is

$$L_c(\theta) = \prod_{i=1}^n f(y_i | z_i; \theta) f(z_i; \theta) = \prod_{i=1}^n [\phi(y_i, v_i^T \beta_1, \sigma_1)^{z_i} \phi(y_i, v_i^T \beta_2, \sigma_2)^{1-z_i} \pi^{z_i} (1 - \pi)^{1-z_i}] ,$$

and its logarithm is

$$\ell_c(\theta) = \sum_{i=1}^n [z_i \log \phi(y_i, v_i^T \beta_1, \sigma_1) + (1 - z_i) \log \phi(y_i, v_i^T \beta_2, \sigma_2) + z_i \log \pi + (1 - z_i) \log(1 - \pi)] .$$

In the E-step, we can compute the conditional expectations  $z_i^{(t)}$  as

$$z_i^{(t)} = \mathbb{E}_{\theta^{(t)}}[Z_i | y_i] = \mathbb{P}_{\theta^{(t)}}[Z_i = 1 | y_i] = \frac{\phi(y_i; v_i^T \beta_1^{(t)}, \sigma_1^{(t)}) \pi^{(t)}}{\phi(y_i; v_i^T \beta_1^{(t)}, \sigma_1^{(t)}) \pi^{(t)} + \phi(y_i; v_i^T \beta_2^{(t)}, \sigma_2^{(t)}) (1 - \pi^{(t)})} \quad (2)$$

and

$$Q(\boldsymbol{\theta}, \boldsymbol{\theta}^{(t)}) = \underbrace{\sum_{i=1}^n z_i^{(t)} \log \phi(y_i, v_i^T \beta_1, \sigma_1)}_A + \underbrace{\sum_{i=1}^n (1 - z_i^{(t)}) \log \phi(y_i, v_i^T \beta_2, \sigma_2)}_B + \underbrace{\log \pi \sum_{i=1}^n z_i^{(t)} + \log(1 - \pi) \left( n - \sum_{i=1}^n z_i^{(t)} \right)}_C.$$

In the M-step, we maximize  $Q(\boldsymbol{\theta}, \boldsymbol{\theta}^{(t)})$  with respect to  $\boldsymbol{\theta}$ . In the above equation,  $A$  depends only on  $\beta_1$  and  $\sigma_1$ ,  $B$  depends only on  $\beta_2$  and  $\sigma_2$  and  $C$  depends on  $\pi$ . We can thus maximize these three terms separately. Reasoning as in Question 2c, we get the following update equations:

$$\begin{aligned} \beta_1^{(t+1)} &= \left[ \sum_{i=1}^n z_i^{(t)} v_i v_i^T \right]^{-1} \sum_{i=1}^n z_i^{(t)} v_i y_i, \\ \sigma_1^{(t+1)} &= \sqrt{\frac{\sum_{i=1}^n z_i^{(t)} (y_i - v_i^T \beta_1^{(t+1)})^2}{\sum_{i=1}^n z_i^{(t)}}}, \\ \beta_2^{(t+1)} &= \left[ \sum_{i=1}^n (1 - z_i^{(t)}) v_i v_i^T \right]^{-1} \sum_{i=1}^n (1 - z_i^{(t)}) v_i y_i, \\ \sigma_2^{(t+1)} &= \sqrt{\frac{\sum_{i=1}^n (1 - z_i^{(t)}) (y_i - v_i^T \beta_2^{(t+1)})^2}{n - \sum_{i=1}^n z_i^{(t)}}}, \\ \pi^{(t+1)} &= \frac{1}{n} \sum_{i=1}^n z_i^{(t)}. \end{aligned}$$

We can now a function that computes the observed-data log-likelihood:

```
loglik_reg1 <- function(theta, y, v) {
  n <- length(y)
  phi1 <- vector(n, mode = "numeric")
  phi2 <- vector(n, mode = "numeric")
  for(i in 1:n) {
    phi1[i] <- dnorm(y[i], mean = theta[1] + theta[2] * v[i], sd = theta[3])
    phi2[i] <- dnorm(y[i], mean = theta[4] + theta[5] * v[i], sd = theta[6])
  }
  logL <- sum(log(theta[7] * phi1 + (1 - theta[7]) * phi2))
  return(logL)
}
```

We then write the EM algorithm, which is similar to those of Exercises 1 and 2:

```
em_outlier_reg1 <- function(y, v, theta0, epsi) {
  go_on <- TRUE
  logL0 <- loglik_reg1(theta0, y, v)
  t <- 0
  n <- length(y)
  phi1 <- vector(n, mode = "numeric")
  phi2 <- vector(n, mode = "numeric")
  print(c(t, logL0))
```

```

while(go_on){
  t<-t+1
  # E-step
  for(i in 1:n){
    phi1[i] <- dnorm(y[i],mean=theta0[1]+theta0[2]*v[i],sd=theta0[3])
    phi2[i] <- dnorm(y[i],mean=theta0[4]+theta0[5]*v[i],sd=theta0[6])
  }
  z<- phi1*theta0[7]/(phi1*theta0[7]+phi2*(1-theta0[7]))
  # M-step
  S<- sum(z)
  pi<-S/n
  reg1<-lm(y ~v,weights=z)
  beta1<-reg1$coefficients
  reg2<-lm(y ~v,weights=1-z)
  beta2<-reg2$coefficients
  sig1<-sqrt(sum(z*reg1$residuals^2)/S)
  sig2<-sqrt(sum((1-z)*reg2$residuals^2)/(n-S))
  theta<-c(beta1,sig1,beta2,sig2,pi)
  logL<-loglik_reg1(theta,y,v)
  if (logL-logL0 < epsi){
    go_on <- FALSE
  }
  logL0 <- logL
  theta0<-theta
  print(c(t,logL))
}
return(list(loglik=logL,theta=theta,z=z))
}

```

### Question 3d

We first initialize the coefficients  $\beta_1$  to the OLS estimates and  $\beta_2$  randomly:

```

beta10 <- reg$coefficients
sig10 <- sd(reg$residuals)
beta20 <- 0.1*rnorm(2)
sig20 <- sig10
pi0 <- 0.8
theta0<-c(beta10,sig10,beta20,sig20,pi0)

```

We then run function `em_outlier_reg1`:

```

estim <- em_outlier_reg1(y,v,theta0,epsi=1e-6)

```

```

## [1] 0.0000 -271.8457
## [1] 1.0000 -244.8924
## [1] 2.0000 -207.9936
## [1] 3.0000 -173.5814
## [1] 4.0000 -196.9353

```

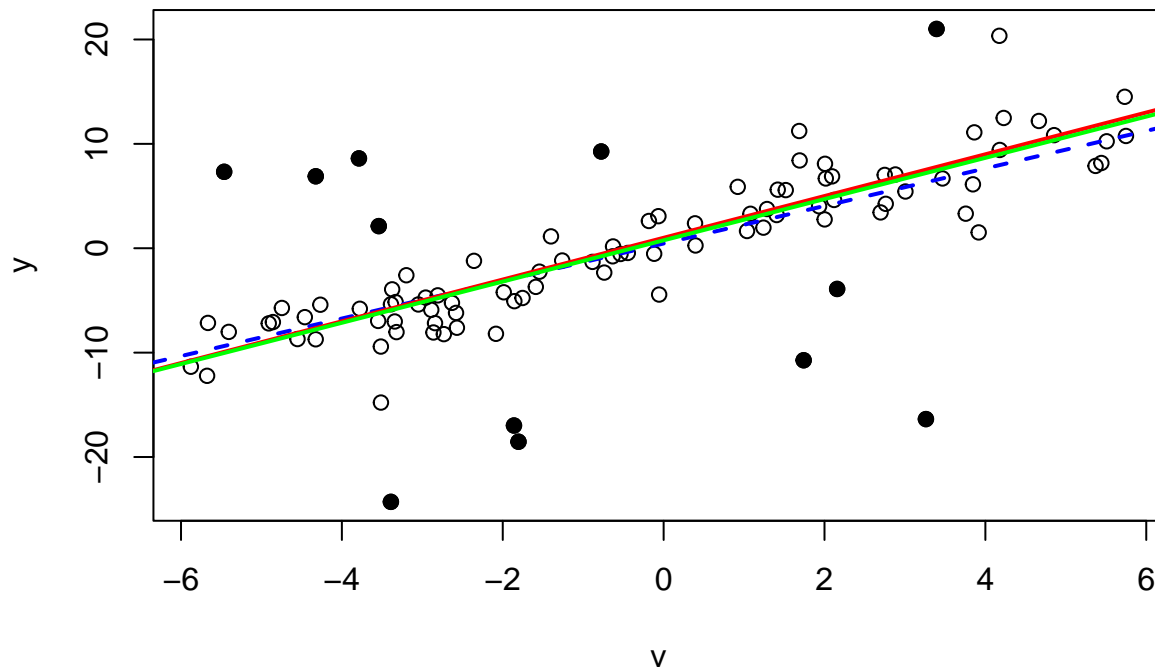
Finally, we plot the line estimated by EM, together with the true regression line and the OLS line. We also plot the points identified as outliers:

```

plot(v,y)
abline(reg,lty=2,col="blue",lwd=2) # LS line

```

```
abline(1,2,lty=1,col="red",lwd=2) # regression line
abline(estim$theta[1],estim$theta[2],lty=1,col='green',lwd=2) # line estimated using EM
points(v[estim$z<0.5],y[estim$z<0.5],pch=19)
```



## Exercise 4

### Question 3a

We write a function `gen_data` that generates a sample:

```
gen_sample<- function(mu,sig,pi,a,n){
  y<-rep(0,n)
  for(i in 1:n){
    z<-sample(c(1,0),size=1,prob=c(pi,1-pi))
    if(z==1) y[i]<- rnorm(1,mean=mu,sd=sig) else y[i]<-runif(1,min=-a,max=a)
  }
  return(y)
}
```

We then define true values of the parameters:

```
pi<-0.90
mu<-0
sig<- 1
theta_true<-c(mu,sig,pi)
```

We then generate  $N = 1000$  samples and estimate  $\theta$  from each sample using function `em_outlier`; the estimates  $\hat{\theta}$  are stored in a matrix `Theta`:

```
a=5
c<-1/(2*a)
n<- 100
```

```

N<-1000
Theta<-matrix(0,N,3)
pi0<-0.5
for(j in 1:N){
  y<- gen_sample(mu,sig,pi,a,n)
  mu0<-mean(y)
  sig0<-sd(y)
  theta0<-c(mu0,sig0,pi0)
  em<-em_outlier(y,theta0,a,eps=1e-5,disp=FALSE)
  Theta[j,<-em$theta }

```

Finally, we can estimate the covariance matrix of  $\hat{\theta}$ :

```

V<-var(Theta)
print(V,2)

```

```

##          [,1]      [,2]      [,3]
## [1,]  1.4e-02 -0.0008 -9.8e-05
## [2,] -8.0e-04  0.0111  2.8e-03
## [3,] -9.8e-05  0.0028  3.0e-03

```

### Question 3b

We first need to compute the complete-data information matrix

$$\hat{i}_{\mathbf{x}}(\hat{\theta}) = \mathbb{E}_{\hat{\theta}}(-\ell''_c(\hat{\theta}) \mid \mathbf{y}).$$

The complete-data likelihood is

$$\begin{aligned}
L_c(\theta) &= \prod_{i=1}^n f(y_i, z_i; \theta) = \prod_{i=1}^n f(y_i \mid z_i; \mu, \sigma) f(z_i; \pi) \\
&= \prod_{i=1}^n [\phi(y_i; \mu, \sigma)^{z_i} c^{1-z_i} \pi^{z_i} (1-\pi)^{1-z_i}]
\end{aligned}$$

and the corresponding log-likelihood is:

$$\ell_c(\theta) = \sum_{i=1}^n z_i \log \phi(y_i; \mu, \sigma) + \left( n - \sum_{i=1}^n z_i \right) \log c + \sum_{i=1}^n (z_i \log \pi + (1 - z_i) \log(1 - \pi))$$

We need to compute the first and then the second derivative. We have

$$\begin{aligned}
\frac{\partial \ell_c}{\partial \mu} &= \frac{1}{\sigma^2} \sum_i z_i (y_i - \mu) \\
\frac{\partial \ell_c}{\partial \sigma} &= -\frac{1}{\sigma} \sum_i z_i + \frac{1}{\sigma^3} \sum_i z_i (y_i - \mu)^2 \\
\frac{\partial \ell_c}{\partial \pi} &= \sum_i \left( \frac{z_i}{\pi} - \frac{1 - z_i}{1 - \pi} \right)
\end{aligned}$$

And then the second derivatives:

$$\frac{\partial^2 \ell_c}{\partial \mu^2} = -\frac{1}{\sigma^2} \sum_i z_i$$

$$\begin{aligned}
\frac{\partial^2 \ell_c}{\partial \mu \partial \sigma} &= -\frac{2}{\sigma^3} \sum_i z_i (y_i - \mu) \\
\frac{\partial^2 \ell_c}{\partial \sigma^2} &= \frac{1}{\sigma^2} \sum_i z_i - \frac{3}{\sigma^4} \sum_i z_i (y_i - \mu)^2 \\
\frac{\partial^2 \ell_c}{\partial \pi^2} &= \sum_i \left( \frac{-z_i}{\pi^2} - \frac{1 - z_i}{(1 - \pi)^2} \right) = -\frac{\sum_i z_i}{\pi^2} - \frac{n - \sum_i z_i}{(1 - \pi)^2} \\
\frac{\partial^2 \ell_c}{\partial \mu \partial \pi} &= \frac{\partial^2 \ell_c}{\partial \sigma \partial \pi} = 0.
\end{aligned}$$

To compute the conditional expectation of  $\ell_c(\boldsymbol{\theta}^{(t)})$  for  $\boldsymbol{\theta} = \boldsymbol{\theta}^{(t)}$ , we simply replace  $z_i$  by  $z_i^{(t)}$  in the above expression. (When the EM algorithm has converged,  $\boldsymbol{\theta}^{(t)} \approx \hat{\boldsymbol{\theta}}$ ). The complete information matrix is then

$$\hat{i}_{\mathbf{x}}(\hat{\boldsymbol{\theta}}) = \begin{pmatrix} \frac{1}{\sigma^2} \sum_i z_i^{(t)} & \frac{2}{\sigma^3} \sum_i z_i^{(t)} (y_i - \hat{\mu}) & 0 \\ \frac{2}{\sigma^3} \sum_i z_i^{(t)} (y_i - \hat{\mu}) & -\frac{1}{\sigma^2} \sum_i z_i^{(t)} + \frac{3}{\sigma^4} \sum_i z_i^{(t)} (y_i - \hat{\mu})^2 & 0 \\ 0 & 0 & \frac{\sum_i z_i^{(t)}}{\pi^2} + \frac{n - \sum_i z_i^{(t)}}{(1 - \pi)^2} \end{pmatrix}.$$

The following R function computes the complete information matrix:

```
complete_information<- function(theta,y,z){
  I<-matrix(0,3,3)
  mu<-theta[1]
  sig<-theta[2]
  pi<-theta[3]
  S<-sum(z)
  I[1,1]<- S/sig^2
  I[1,2]<- 2/sig^3*sum(z*(y-mu))
  I[2,1]<-I[1,2]
  I[2,2]<- -S/sig^2 + 3/sig^4*sum(z*(y-mu)^2)
  I[3,3]<- S/pi^2 + (n-S)/(1-pi)^2
  return(I)
}
```

Next, we need to compute the missing information as the variance of the conditional score:

$$\hat{i}_{\mathbf{Z}|\mathbf{Y}}(\boldsymbol{\theta}) = \text{Var} \left[ \frac{\partial \log f(\mathbf{z} | \mathbf{y}; \boldsymbol{\theta})}{\partial \boldsymbol{\theta}} \mid \mathbf{y} \right].$$

For this, we first need to compute  $f(\mathbf{z} | \mathbf{y}; \boldsymbol{\theta})$ . We recall that

$$P(Z_i = 1 \mid y_i) = \frac{\phi_i \pi}{\phi_i \pi + c(1 - \pi)} = p_i(\boldsymbol{\theta}).$$

with

$$\phi_i = \phi(y_i; \mu, \sigma) = \frac{1}{\sigma \sqrt{2\pi}} \exp \left[ -\frac{1}{2\sigma^2} (y_i - \mu)^2 \right].$$

So,

$$f(\mathbf{z} | \mathbf{y}; \boldsymbol{\theta}) = \prod_{i=1}^n f(z_i \mid y_i; \boldsymbol{\theta}) = \prod_{i=1}^n p_i(\boldsymbol{\theta})^{z_i} (1 - p_i(\boldsymbol{\theta}))^{1-z_i},$$

and

$$\begin{aligned}
\log f(\mathbf{z} | \mathbf{y}; \boldsymbol{\theta}) &= \sum_{i=1}^n (z_i (\log \phi_i + \log \pi - \log[\phi_i \pi + c(1 - \pi)]) + (1 - z_i) (\log c(1 - \pi) - \log[\phi_i \pi + c(1 - \pi)])) \\
&= \sum_{i=1}^n (z_i \log \phi_i + z_i \log \pi + (1 - z_i) \log c(1 - \pi) - \log[\phi_i \pi + c(1 - \pi)]).
\end{aligned}$$



To compute the score, we first compute:

$$\frac{\partial \phi_i}{\partial \mu} = \phi_i \frac{y_i - \mu}{\sigma^2},$$

and

$$\frac{\partial \phi_i}{\partial \sigma} = \frac{\phi_i}{\sigma} \left( \frac{(y_i - \mu)^2}{\sigma^2} - 1 \right).$$

Using elementary calculus, we get

$$\begin{aligned} \frac{\partial \log f(\mathbf{z} | \mathbf{y}; \boldsymbol{\theta})}{\partial \mu} &= \frac{1}{\sigma^2} \sum_{i=1}^n z_i (y_i - \mu) - \frac{\pi}{\sigma^2} \sum_{i=1}^n \frac{(y_i - \mu) \phi_i}{\phi_i \pi + c(1 - \pi)} \\ \frac{\partial \log f(\mathbf{z} | \mathbf{y}; \boldsymbol{\theta})}{\partial \sigma} &= \frac{1}{\sigma} \sum_{i=1}^n z_i \left( \frac{(y_i - \mu)^2}{\sigma^2} - 1 \right) - \frac{\pi}{\sigma} \sum_{i=1}^n \frac{\phi_i \left( \frac{(y_i - \mu)^2}{\sigma^2} - 1 \right)}{\phi_i \pi + c(1 - \pi)} \\ \frac{\partial \log f(\mathbf{z} | \mathbf{y}; \boldsymbol{\theta})}{\partial \pi} &= \frac{\sum_{i=1}^n z_i - n\pi}{\pi(1 - \pi)} - \sum_{i=1}^n \frac{\phi_i - c}{\phi_i \pi + c(1 - \pi)}. \end{aligned}$$

The calculation of the score is performed by the following R function:

```
score<- function(theta,y,z,c){
  S<-c(0,0,0)
  mu<-theta[1]
  sig<-theta[2]
  pi<-theta[3]
  SZ<-sum(z)
  phi<- sapply(y,dnorm,mean=mu,sd=sig)
  S[1]<- 1/sig^2*sum(z*(y-mu)) - pi/sig^2*sum((y-mu)*phi/(phi*pi+c*(1-pi)))
  S[2]<- -SZ/sig + 1/sig^3*sum(z*(y-mu)^2) -
    pi/sig*sum( phi*((y-mu)^2/sig^2-1)/(phi*pi+c*(1-pi)) )
  S[3]<- (SZ-n*pi)/(pi*(1-pi)) - sum((phi-c)/(phi*pi+c*(1-pi)))
  return(S)
}
```

We can now compute the observed information for an observed sample. We first generate the data and compute the MLE:

```
set.seed(42)
y<- gen_sample(mu,sig,pi,a,n)
mu0<-mean(y)
sig0<-sd(y)
pi0<-0.5
theta0<-c(mu0,sig0,pi0)
em<-em_outlier(y,theta0,a,epsi=1e-5)
```

```
## [1] 0.0000 -191.3151
## [1] 1.0000 -173.8608
## [1] 2.0000 -169.5465
## [1] 3.0000 -168.7118
## [1] 4.0000 -168.4584
## [1] 5.0000 -168.3611
## [1] 6.0000 -168.3219
## [1] 7.0000 -168.3059
## [1] 8.0000 -168.2994
## [1] 9.0000 -168.2968
## [1] 10.0000 -168.2957
```

```
## [1] 11.0000 -168.2953
## [1] 12.0000 -168.2951
## [1] 13.0000 -168.295
## [1] 14.0000 -168.295
## [1] 15.0000 -168.295
## [1] 16.0000 -168.295
```

We then compute by simulation the missing information  $\hat{i}_{\mathbf{Z}|\mathbf{Y}}(\boldsymbol{\theta})$  as the variance of the score:

```
N<-100000
SS<-matrix(0,N,3)
for(i in (1:N)){
  u<-runif(n)
  Z<-as.numeric(u<=em$z)
  SS[i,]<-score(em$theta,y,Z,c)
}
izy<-var(SS)
```

Finally, we compute the observed information as the difference between the complete information and missing information; and we estimate the variance of  $\hat{\boldsymbol{\theta}}$  as the inverse of the observed information:

```
ix<-complete_information(em$theta,y,em$z)
iy<-ix-izy
varh<-solve(iy)
print(varh,2)
```

```
##          [,1]      [,2]      [,3]
## [1,]  0.01254 -0.0011 -0.00047
## [2,] -0.00111  0.0086  0.00175
## [3,] -0.00047  0.0018  0.00315
```

To check our calculations, we can compare our result to that using numerical differentiation:

```
library(numDeriv)
varh1<-solve(-hessian(loglik,em$theta,y=y))
print(varh1,2)
```

```
##          [,1]      [,2]      [,3]
## [1,]  0.01254 -0.0011 -0.00046
## [2,] -0.00112  0.0087  0.00179
## [3,] -0.00046  0.0018  0.00318
```