

# Incomplete Data Analysis

## Assignment 2

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### Question 1

Suppose  $Y_1, \dots, Y_n$  are independent and identically distributed with cumulative distribution function given by

$$F(y; \theta) = 1 - e^{-y^2/(2\theta)}, \quad y \geq 0, \theta > 0$$

Further suppose that observations are (right) censored if  $Y_i > C$ , for some known  $C > 0$ , and let

$$X_i = \begin{cases} Y_i & \text{if } Y_i \leq C, \\ C & \text{if } Y_i > C, \end{cases} \quad R_i = \begin{cases} 1 & \text{if } Y_i \leq C \\ 0 & \text{if } Y_i > C \end{cases}$$

### Question 1a

Show that the maximum likelihood estimator based on the observed data  $\{x_i, r_i\}_{i=1}^n$  is given by

$$\hat{\theta} = \frac{\sum_{i=1}^n X_i^2}{2 \sum_{i=1}^n R_i}$$

### Solution:

- To derive the MLE we must maximize the log-likelihood of the observed data  $\{(x_i, r_i)\}_{i=1}^n$ . In this context, there are two contributions to the likelihood function:
  1.  $f(y_i; \theta) = \frac{dF(y_i; \theta)}{dy_i}$  from *non-censored* observations.
  2.  $Pr(Y_i > C; \theta) = S(C; \theta) = 1 - F(y_i; \theta)$  from *censored* observations.
- All observations  $Y_i, \dots, Y_n$  are iid, hence,

$$\begin{aligned} L(\theta) &= \prod_{i=1}^n \{ [f(y_i; \theta)]^{r_i} [S(C; \theta)]^{1-r_i} \} \\ &= \prod_{i=1}^n \left\{ \left[ \frac{y_i}{\theta} e^{-y_i^2/2\theta} \right]^{r_i} \left[ e^{-C^2/2\theta} \right]^{1-r_i} \right\} \\ &= \left( \frac{y_i}{2\theta} \right)^{\sum_i r_i} \exp \left( -\frac{1}{2\theta} \sum_i [r_i y_i^2 + (1-r_i)C^2] \right) \\ &= \left( \frac{y_i}{2\theta} \right)^{\sum_i r_i} \exp \left( -\frac{1}{2\theta} \sum_i x_i^2 \right) \end{aligned}$$

- Note that in order to understand how one goes from line 3 to line 4 in the equation defined above, we recall that we can write the variable  $X_i$  as  $X_i = Y_i R_i + C(1 - R_i)$  and due to the binary nature of  $R_i$ :

$$\begin{aligned}\implies X_i^2 &= Y_i^2 R_i^2 + C^2(1 - R_i)^2 + 2Y_i R_i C(1 - R_i) \\ \implies X_i^2 &= Y_i^2 R_i + C^2(1 - R_i)\end{aligned}$$

- We can now define the log-likelihood to be

$$\log L(\theta) := l(\theta) = \sum_{i=1}^n r_i \log\left(\frac{y_i}{2\theta}\right) - \frac{1}{2\theta} \sum_{i=1}^n x_i^2$$

- Maximising this quantity through taking its derivative

$$\frac{d}{d\theta} l(\theta) = -\frac{\sum_{i=1}^n r_i}{\theta} + \frac{\sum_{i=1}^n x_i^2}{2\theta^2}$$

- leading to

$$\hat{\theta}_{\text{MLE}} = \frac{\sum_{i=1}^n X_i^2}{2 \sum_{i=1}^n R_i}.$$

- Note that we have assumed here that  $\hat{\theta}_{\text{MLE}}$  is indeed a maximum and have not computed the second derivative since our result matches the one given in the question.

### Question 1b

Show that the expected Fisher information for the observed data likelihood is

$$I(\theta) = \frac{n}{\theta^2} (1 - e^{-C^2/2\theta})$$

**Note:**  $\int_0^C y^2 f(y; \theta) dy = -C^2 e^{-C^2/2\theta} + 2\theta(1 - e^{-C^2/2\theta})$ , where  $f(y; \theta)$  is the density function corresponding to the cumulative distribution function  $F(y; \theta)$  defined above.

### Solution:

- We first recall the general definition of the expected Fisher information to be

$$I(\theta) = -\mathbb{E} \left[ \frac{d^2 l(\theta)}{d\theta^2} \right]$$

- We now compute the second derivative of the log-likelihood and re-introduce the variables  $r_i$  and  $y_i$  for  $x_i$  which will allow us to take expectations more clearly. This yields,

$$I(\theta) = -\frac{\sum_i \mathbb{E}[R_i]}{\theta^2} + \frac{\sum_i \mathbb{E}[R_i Y_i^2]}{\theta^3} + \frac{\sum_i C^2 \mathbb{E}[(1 - R_i)]}{\theta^3}$$

- Note that  $R$  is a binary random variable and so

$$\begin{aligned}
\mathbb{E}[R] &= 1 \times \Pr(R = 1) + 0 \times \Pr(R = 0) \\
&= \Pr(R = 1) \\
&= \Pr(Y \leq C) \\
&= F(C; \theta) \\
&= 1 - e^{-C^2/2\theta}.
\end{aligned}$$

- And hence

$$\begin{aligned}
I(\theta) &= -\frac{\sum_i \mathbb{E}[R_i]}{\theta^2} + \frac{\sum_i \mathbb{E}[R_i Y_i^2]}{\theta^3} + \frac{\sum_i C^2 \mathbb{E}[(1 - R_i)]}{\theta^3} \\
&= -\frac{n}{\theta^2}(1 - e^{-C^2/2\theta}) + \frac{n}{\theta^3} \left\{ -C^2 e^{-C^2/2\theta} + 2\theta(1 - e^{-C^2/2\theta}) \right\} + \frac{n}{\theta^3} e^{-C^2/2\theta} \\
&= \frac{n}{\theta^2}(1 - e^{-C^2/2\theta})
\end{aligned}$$

- Where the note given in the question was used to calculate  $\mathbb{E}[R_i Y_i^2] = \int_0^C y^2 f(y; \theta) dy = -C^2 e^{-C^2/2\theta} + 2\theta(1 - e^{-C^2/2\theta})$

### Question 1c

Appealing to the asymptotic normality of the maximum likelihood estimator, provide a 95% confidence interval for  $\theta$ .

**Solution:**

- We recall the asymptotic normality of the MLE as

$$\hat{\theta}_{\text{MLE}} \sim N(\theta, I(\theta)^{-1})$$

- Therefore

$$\frac{\hat{\theta}_{\text{MLE}} - \theta}{\sqrt{I(\theta)^{-1}}} \sim N(0, 1)$$

- Using the properties of the standard Gaussian distribution ( $\alpha = 0.05$ )

$$\Pr \left( z_{-\alpha/2} \leq \frac{\hat{\theta}_{\text{MLE}} - \theta}{\sqrt{I(\theta)^{-1}}} \leq z_{\alpha/2} \right) = 1 - \alpha = 0.95$$

- The 95% CI for  $\theta$  is hence  $\left[ \sqrt{I(\theta)^{-1}} z_{-\alpha/2} + \hat{\theta}_{\text{MLE}}, \sqrt{I(\theta)^{-1}} z_{\alpha/2} + \hat{\theta}_{\text{MLE}} \right]$  where  $z_{\alpha/2} = 1.959964$ ,  $z_{-\alpha/2} = -1.959964$ , and  $\sqrt{I(\theta)^{-1}} = \theta / \sqrt{n(1 - e^{-C^2/2\theta})}$

```
alpha = 0.05
```

```
z = qnorm(1-alpha/2)
```

## Question 2

Suppose that  $Y_i \sim N(\mu, \sigma^2)$  are iid for  $i = 1, \dots, n$ . Further suppose that now observations are (left) censored if  $Y_i < D$ , for some known  $D$  and let

$$X_i = \begin{cases} Y_i & \text{if } Y_i \geq D, \\ D & \text{if } Y_i < D, \end{cases} \quad R_i = \begin{cases} 1 & \text{if } Y_i \geq D \\ 0 & \text{if } Y_i < D \end{cases}$$

### Question 2a

Show that the log-likelihood of the observed data  $\{x_i, r_i\}_{i=1}^n$  is given by

$$l(\mu, \sigma^2 | \mathbf{x}, \mathbf{r}) = \sum_{i=1}^n \{r_i \log \phi(x_i; \mu, \sigma^2) + (1 - r_i) \log \Phi(x_i; \mu, \sigma^2)\}$$

where  $\phi(x_i; \mu, \sigma^2)$  and  $\Phi(x_i; \mu, \sigma^2)$  stands, respectively, for the density function and cumulative distribution function of the normal distribution with mean  $\mu$  and variance  $\sigma^2$ .

**Solution:**

- Similar to 1a, our likelihood function has two contributions:
  1.  $\phi(x_i; \mu, \sigma^2)$  from *non-censored* observations.
  2.  $Pr(X_i < D; \mu, \sigma^2) = S(D; \mu, \sigma^2) = \Phi(x_i; \mu, \sigma^2)$  from *censored* observations.
- All observations  $X_i, \dots, X_n$  are iid, hence,

$$\begin{aligned} l(\mu, \sigma^2 | \mathbf{x}, \mathbf{r}) &= \log \prod_{i=1}^n \{ \phi(x_i; \mu, \sigma^2)^{r_i} [\Phi(x_i; \mu, \sigma^2)]^{1-r_i} \} \\ &= \log \left\{ \phi(x_i; \mu, \sigma^2)^{\sum_i r_i} [\Phi(x_i; \mu, \sigma^2)]^{\sum_i (1-r_i)} \right\} \\ &= \sum_{i=1}^n \{ r_i \log \phi(x_i; \mu, \sigma^2) + (1 - r_i) \log \Phi(x_i; \mu, \sigma^2) \} \end{aligned}$$

### Question 2b

Determine the maximum likelihood estimate of  $\mu$  based on the data available in the file `dataex2.Rdata`. Consider  $\sigma^2$  known and equal to  $1.5^2$ .

**Solution:**

- Given that we are now in possession of the log-likelihood of our data, we proceed to maximize this with respect to the parameter  $\mu$  in order to derive our MLE of this parameter which we denote  $\hat{\mu}_{\text{MLE}}$ .
- The Newton-Raphson method was used to optimise our log-likelihood which yielded the following MLE:  $\hat{\mu}_{\text{MLE}} = 5.5328$  to 4 d.p.

```
library(maxLik)

# Loading in data

load('dataex2.Rdata')
```

```

# Log likelihood function set to maximized
get_log_likelihood = function(param, data) {
  mu = param
  x = data[,1]; r = data[,2]
  return(sum(r*dnorm(x, mean=mu, sd=1.5, log=TRUE) +
            (1 - r)*pnorm(x, mean=mu, sd=1.5, log.p=TRUE)))
}

# Get MLE
mle = maxLik(logLik = get_log_likelihood, data = dataex2, start = c(mu=1))

# Present results
summary(mle)

```

```

## -----
## Maximum Likelihood estimation
## Newton-Raphson maximisation, 3 iterations
## Return code 2: successive function values within tolerance limit
## Log-Likelihood: -336.3821
## 1 free parameters
## Estimates:
##      Estimate Std. error t value Pr(> t)
## mu    5.5328      0.1075   51.48 <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## -----

```

### Question 3

Consider a bivariate normal sample  $(Y_1, Y_2)$  with parameters  $\theta = (\mu_1, \mu_2, \sigma_1^2, \sigma_{12}, \sigma_2^2)$ . The variable  $Y_1$  is fully observed, while some values of  $Y_2$  are missing. Let  $R$  be the missingness indicator, taking the value 1 for observed values and 0 for missing values. For the following missing data mechanisms state, justifying, whether they are ignorable for likelihood-based estimation.

#### Solution:

- A missing data mechanism (MDM) is said to be ignorable for likelihood based inference if and only if the following two criteria are met:
  1. The missing data are missing at random (MAR) or missing completely at random (MCAR).
  2. The parameter  $\psi$  (missingness mechanism) and  $\theta$  (data model) are distinct in the sense that the joint parameter space of  $(\psi, \theta)$  is the product of the parameter spaces  $\Psi$  and  $\Theta$  (separability condition).

(a)  $\text{logit}\{Pr(R=0|y_1, y_2, \theta, \psi)\} = \psi_0 + \psi_1 y_1$ ;  $\psi = (\psi_1, \psi_2)$  distinct from  $\theta$ .

- We observe that the MDM is dependent on the fully observed variable,  $y_1$ , only. The missing data resulting from this mechanism will hence be MAR. Furthermore,  $\psi$  (missingness mechanism) and  $\theta$  (data model) are distinct indicating that MDM (a) is ignorable for likelihood based inference.

(b)  $\text{logit}\{Pr(R=0|y_1, y_2, \theta, \psi)\} = \psi_0 + \psi_1 y_2$ ;  $\psi = (\psi_1, \psi_2)$  distinct from  $\theta$ .

- We observe that the MDM is dependent on the missing variable,  $y_2$ , only. The missing data resulting from this mechanism will hence be missing not at random (MNAR) indicating that MDM (b) is **NOT** ignorable for likelihood-based inference.

(c)  $\text{logit}\{Pr(R=0|y_1, y_2, \theta, \psi)\} = 0.5(\mu_1 + \psi_1 y_1)$ ;  $\psi$  (scalar) distinct from  $\theta$ .

- We observe a similar MDM to (a) with an added dependency on  $\mu_1$ . The parameter  $\mu_1$  is contained within the parameter space of both the missing data model,  $\theta$ , and the MDM,  $\psi$ . Criterion 2 is hence not met and MDM (c) is **NOT** ignorable for likelihood-based inference.

## Question 4

Suppose that

$$Y_i \stackrel{\text{ind}}{\sim} \text{Bernoulli}(p_i(\boldsymbol{\beta})),$$

$$p_i(\boldsymbol{\beta}) = \frac{\exp(\beta_0 + x_i\beta_1)}{1 + \exp(\beta_0 + x_i\beta_1)}$$

for  $i = 1, \dots, n$  and  $\boldsymbol{\beta} = (\beta_0, \beta_1)'$ . Although the covariate  $x$  is fully observed, the response variable  $Y$  has missing values. Assuming ignorability, derive and implement the EM algorithm to compute the MLE of  $\boldsymbol{\beta}$  based on the data available in `dataex4.Rdata`.

**Solution:**

- We begin by first obtaining the likelihood for  $\boldsymbol{\beta}$  which is given by

$$L(\boldsymbol{\beta}) = \prod_{i=1}^n \{p_i(\boldsymbol{\beta})^{y_i} [1 - p_i(\boldsymbol{\beta})]^{1-y_i}\}$$

$$= \prod_{i=1}^n \left\{ \left( \frac{e^{\beta_0 + x_i\beta_1}}{1 + e^{\beta_0 + x_i\beta_1}} \right)^{y_i} \left( \frac{1}{1 + e^{\beta_0 + x_i\beta_1}} \right)^{1-y_i} \right\}.$$

- The corresponding log likelihood is hence

$$l(\boldsymbol{\beta}) = \sum_{i=1}^n \left\{ y_i \log \left( \frac{e^{\beta_0 + x_i\beta_1}}{1 + e^{\beta_0 + x_i\beta_1}} \right) + (1 - y_i) \log \left( \frac{1}{1 + e^{\beta_0 + x_i\beta_1}} \right) \right\}$$

$$= \sum_{i=1}^n \{y_i(\beta_0 + x_i\beta_1) - \log(1 + e^{\beta_0 + x_i\beta_1})\}.$$

- Now that we are in possession of the log-likelihood, we can use this to conduct the expectation step of the EM algorithm and define our  $Q$  function.
- Note that the expectation is taken under the distribution of the **missing data**. We hence make use of our univariate pattern of missingness and assume that the first  $m$  values of  $Y$  are reserved and the remaining  $n - m$  are missing i.e.  $\mathbf{y}_{obs} = y_1, \dots, y_m$  and  $\mathbf{y}_{mis} = y_{m+1}, \dots, y_n$ .

$$Q(\boldsymbol{\beta}|\boldsymbol{\beta}^{(t)}) = \mathbb{E}_{\mathbf{y}_{mis}} [l(\boldsymbol{\beta})|\mathbf{y}_{obs}, \mathbf{x}, \boldsymbol{\beta}^{(t)}]$$

$$= \sum_{i=1}^m \{y_i(\beta_0 + \beta_1 x_i)\} - \sum_{i=1}^n \log(1 + e^{\beta_0 + \beta_1 x_i}) + \sum_{i=m+1}^n (\beta_0 + \beta_1 x_i) \mathbb{E}_{\mathbf{y}_{mis}} [y_i|\mathbf{y}_{obs}, \mathbf{x}, \boldsymbol{\beta}^{(t)}]$$

$$= \sum_{i=1}^m \{y_i(\beta_0 + \beta_1 x_i)\} - \sum_{i=1}^n \log(1 + e^{\beta_0 + \beta_1 x_i}) + \sum_{i=m+1}^n (\beta_0 + \beta_1 x_i) p_i(\boldsymbol{\beta})$$

- Where we have used the result that  $\mathbb{E}[Y_i] = p_i(\boldsymbol{\beta})$  since  $Y_i \stackrel{\text{ind}}{\sim} \text{Bernoulli}\{p_i(\boldsymbol{\beta})\}$ . We remind the reader of the definition of  $p_i(\boldsymbol{\beta})$  is  $p_i(\boldsymbol{\beta}) = \frac{\exp(\beta_0 + x_i\beta_1)}{1 + \exp(\beta_0 + x_i\beta_1)}$  as defined in the question.
- Following our definition of the  $Q$  function, we conduct the maximization step of the EM algorithm and maximize this function with respect to the parameters,  $\boldsymbol{\beta} = (\beta_0, \beta_1)'$ .
- The R code below repeatedly maximizes our  $Q$  function for every iteration of our algorithm in order to find subsequent values of the parameters considered.
- This process is repeated until the following convergence criterion is met and we have our MLEs ( $\epsilon = 1 \times 10^{-10}$ ):

$$|p^{(t+1)} - p^{(t)}| + |\mu^{(t+1)} - \mu^{(t)}| + |(\sigma^{(t+1)})^2 - (\sigma^{(t)})^2| + |\lambda^{(t+1)} - \lambda^{(t)}| < \epsilon$$

- At convergence, the following results are achieved for our MLEs:  $\widehat{\beta}_{0\text{MLE}} = 0.9755$  to 4 d.p and  $\widehat{\beta}_{1\text{MLE}} = -2.4804$  to 4 d.p as our MLEs for  $\beta = (\beta_0, \beta_1)'$ .

```
# Loading relevant packages
library(maxLik)
library(dplyr)
library(tidyr)
library(magrittr)

# Loading data
load('dataex4.Rdata')

dataex4 = dataex4 %>%
  # Sorting data in ascending order in column Y (0 -> 1 -> NA)
  arrange(Y) %>%
  # Creating indicator variable column (if Y == NA -> R = 0, else -> R = 1)
  mutate(R = (Y == 0 | Y == 1)*1) %>%
  # Replacing NAs with 0s in R column
  tidyr::replace_na(list(R = 0)) %>%
  # Replacing NAs with 2s in Y column to prevent coercion problems i.e. 0*NA=NA
  tidyr::replace_na(list(Y = 2))

# Sigmoid probability function
prob = function(beta, x) {
  return(exp(beta[1] + x*beta[2]) / (1 + exp(beta[1] + x*beta[2])))
}

# Defining Q function for EM algorithm
q_function = function(params, data){
  beta0 = params[1]; beta1 = params[2]
  xx = data$X
  yy = data$Y
```



```

rr = data$R
sum(yy*rr*(beta0 + beta1*xx) - log(1 + exp(beta0 + beta1*xx)) +
    (1 - rr)*(beta0 + beta1*xx)*prob(beta.old, xx))
}

# Repeatedly maximising to get \beta^{(t+1)}
# until convergence criterion is met
tol = 1e-10
beta.old = c(0, 0)
repeat{
  beta = coef(maxLik(q_function, data=dataex4, start = beta.old))
  if (max(abs(beta - beta.old)) < tol) {
    break
  }
  beta.old = beta
}
beta

```

```
## [1] 0.9755261 -2.4803837
```

### Question 5

Consider a random sample  $Y_1, \dots, Y_n$  from the mixture distribution with density

$$f(y) = pf_{\log\text{Normal}}(y; \mu, \sigma^2) + (1 - p)f_{\text{Exp}}(y; \lambda),$$

with

$$f_{\log\text{Normal}}(y; \mu, \sigma^2) = \frac{1}{y\sqrt{2\pi\sigma^2}} \exp\left\{\frac{1}{2\sigma^2}(\log y - \mu)^2\right\}, \quad y > 0, \mu \in \mathbb{R}, \sigma > 0$$

$$f_{\text{Exp}}(y; \lambda) = \lambda e^{-\lambda y}, \quad y \geq 0, \lambda > 0$$

and  $\theta = (p, \mu, \sigma^2, \lambda)$

### Question 5a

Derive the EM algorithm to find the updating equations for  $\theta^{(t+1)} = (p^{(t+1)}, \mu^{(t+1)}, (\sigma^{(t+1)})^2, \lambda^{(t+1)})$ .

**Solution:**

- We begin by taking our mixture model and computing the likelihood:

$$L(\theta; y) = \prod_{i=1}^n \{pf_{\log\text{Normal}}(y_i; \mu, \sigma^2) + (1 - p)f_{\text{Exp}}(y_i; \lambda)\}$$

- The log likelihood is hence

$$l(\theta; y) = \sum_{i=1}^n \log \{pf_{\log\text{Normal}}(y_i; \mu, \sigma^2) + (1 - p)f_{\text{Exp}}(y_i; \lambda)\}$$

- The combination of the summation and logarithm make it very difficult to maximize this log-likelihood. We hence turn to the EM algorithm for this computation, but first, we must artificially create *missing data* in order to implement it.
- The idea that enables us to implement the EM algorithm is that if we knew the group the observation  $y_i$  belonged to, then we could fit the appropriate distribution (log-normal or exponential).
- Let us define  $\mathbf{y}_{\text{obs}} = (y_1, \dots, y_n)$  and  $\mathbf{y}_{\text{mis}} = \mathbf{z} = (z_1, \dots, z_n)$  where  $Z_i \sim \text{Bernoulli}(p)$  and hence

$$Z_i = \begin{cases} 1 & \text{if } Y_i \text{ belongs to } f_{\log\text{Normal}}(y_i; \mu, \sigma^2) \\ 0 & \text{if } Y_i \text{ belongs to } f_{\text{Exp}}(y_i; \lambda) \end{cases}$$

- Re-writing our log-likelihood in terms of  $\mathbf{z}$

$$l(\theta; y, \mathbf{z}) = \sum_{i=1}^n z_i \{\log p + \log f_{\log\text{Normal}}(y_i; \mu, \sigma^2)\} + \sum_{i=1}^n (1 - z_i) \{\log(1 - p) + \log f_{\text{Exp}}(y_i; \lambda)\}$$

- With our log-likelihood and artificial missing data in hand we can now define our Q function to be:

$$Q(\theta|\theta^{(t)}) = \mathbb{E}_{\mathbf{z}} [l(\theta; y, \mathbf{z})]$$

$$= \sum_{i=1}^n \mathbb{E}[\mathbf{z}|\mathbf{y}, \theta^{(t)}] \{\log p + \log f_{\log\text{Normal}}(y_i; \mu, \sigma^2)\} + \sum_{i=1}^n (1 - \mathbb{E}[\mathbf{z}|\mathbf{y}, \theta^{(t)}]) \{\log(1 - p) + \log f_{\text{Exp}}(y_i; \lambda)\}$$

- Taking the relevant expectation

$$\begin{aligned}
\mathbb{E}[z_i | \mathbf{y}, \boldsymbol{\theta}^{(t)}] &= \mathbb{E}[z_i | \mathbf{y}, \boldsymbol{\theta}^{(t)}] \\
&= \Pr(z_i = 1 | y_i, \boldsymbol{\theta}^{(t)}) \\
&= \frac{p_i^{(t)} f_{\log\text{Normal}}(y_i; \mu, \sigma^2)}{p_i^{(t)} f_{\log\text{Normal}}(y_i; \mu, \sigma^2) + (1 - p_i^{(t)}) f_{\text{Exp}}(y_i; \lambda)} \\
&:= \tilde{p}_i^{(t)}
\end{aligned}$$

- We hence define our Q function to be

$$Q(\boldsymbol{\theta} | \boldsymbol{\theta}^{(t)}) = \sum_{i=1}^n \tilde{p}_i^{(t)} \{ \log p + \log f_{\log\text{Normal}}(y_i; \mu, \sigma^2) \} + \sum_{i=1}^n (1 - \tilde{p}_i^{(t)}) \{ \log(1 - p) + \log f_{\text{Exp}}(y_i; \lambda) \}$$

- With the Q function defined above, we can analytically maximize this function in order to find  $\boldsymbol{\theta}^{(t+1)}$ . Taking the derviative of the Q function with respect to all our parameters yields

$$\begin{aligned}
\frac{\partial Q(\boldsymbol{\theta} | \boldsymbol{\theta}^{(t)})}{\partial p} &= \frac{1}{p} \sum_{i=1}^n \tilde{p}_i^{(t)} - \frac{1}{(1-p)} \sum_{i=1}^n (1 - \tilde{p}_i^{(t)}) \\
\frac{\partial Q(\boldsymbol{\theta} | \boldsymbol{\theta}^{(t)})}{\partial \mu} &= \sum_{i=1}^n \tilde{p}_i^{(t)} \left\{ \frac{\log y_i - \mu}{\sigma^2} \right\} \\
\frac{\partial Q(\boldsymbol{\theta} | \boldsymbol{\theta}^{(t)})}{\partial \sigma^2} &= \sum_{i=1}^n \tilde{p}_i^{(t)} \left\{ \frac{(\log y_i - \mu)^2}{2\sigma^4} - \frac{1}{2\sigma^2} \right\} \\
\frac{\partial Q(\boldsymbol{\theta} | \boldsymbol{\theta}^{(t)})}{\partial \lambda} &= \sum_{i=1}^n (1 - \tilde{p}_i^{(t)}) \left\{ \frac{1}{\lambda} - y_i \right\}
\end{aligned}$$

- Setting the each derivative defined above to zero and solving for each parameter in  $\boldsymbol{\theta}$  at iteration  $t+1$  we yield the following update equations

$$\begin{aligned}
p^{(t+1)} &= \frac{1}{n} \sum_{i=1}^n \tilde{p}_i^{(t)} \\
\mu^{(t+1)} &= \frac{\sum_{i=1}^n \tilde{p}_i^{(t)} \log y_i}{\sum_{i=1}^n \tilde{p}_i^{(t)}} \\
(\sigma^{(t+1)})^2 &= \frac{\sum_{i=1}^n \tilde{p}_i^{(t)} (\log y_i - \mu^{(t+1)})^2}{\sum_{i=1}^n \tilde{p}_i^{(t)}} \\
\lambda^{(t+1)} &= \frac{\sum_{i=1}^n (1 - \tilde{p}_i^{(t)})}{\sum_{i=1}^n (1 - \tilde{p}_i^{(t)}) y_i}
\end{aligned}$$

## Question 5b

Using the dataset `datasetex5.Rdata` implement the EM algorithm and find the MLEs for each component of  $\boldsymbol{\theta}$ . As starting values, you might want to consider  $\boldsymbol{\theta}^{(0)} = (p^{(0)}, \mu^{(0)}, (\sigma^{(0)})^2, \lambda^{(0)}) = (0.1, 1, 0.5^2, 2)$ . Draw the histogram of the data with the estimated density superimposed.

### Solution:

- Using the code defined below (see comments for details on implementation), we find the MLEs for each component of  $\boldsymbol{\theta}$  to be (4 d.p):  $\hat{p}_{\text{MLE}} = 0.4795$ ;  $\hat{\mu}_{\text{MLE}} = 2.0133$ ;  $\hat{\sigma}_{\text{MLE}}^2 = 0.8637$ ;  $\hat{\lambda}_{\text{MLE}} = 1.0330$

```

load('dataex5.Rdata')

# Calculates tilde p for all i, with the current parameter values
p_tilde <- function(y, p.t, mu.t, sigma2.t, lambda.t) {
  c1 <- p.t * dlnorm(y, mu.t, sqrt(sigma2.t))
  c2 <- (1 - p.t) * dexp(y, lambda.t)
  c1 / (c1 + c2)
}

# Applies the EM algorithm to fit the mixture density with provided observations y
em <- function(y, p.t, mu.t, sigma2.t, lambda.t, eps = 1e-8, maxit = 1e3) {
  n <- length(y)
  t <- 0

  # Until converged or maximum iterations reached, iterate indefinitely...
  repeat {
    # Calculate the tilde p values for all i, using the current parameter values
    p_tilde.t <- p_tilde(y, p.t, mu.t, sigma2.t, lambda.t)

    # Store tilde p sum for efficiency, since this is used in multiple update equations
    sum.p_tilde.t <- sum(p_tilde.t)

    # Store the previous parameter values (used later for convergence check)
    params.prev <- c(p.t, mu.t, sigma2.t, lambda.t)

    # Update parameters using the update equations
    p.t <- sum.p_tilde.t / n
    mu.t <- sum(p_tilde.t * log(y)) / sum.p_tilde.t
    sigma2.t <- sum(p_tilde.t * (log(y) - mu.t)^2) / sum.p_tilde.t
    lambda.t <- (n - sum.p_tilde.t) / sum((1 - p_tilde.t) * y)

    # Increment iteration counter
    t <- t + 1
  }
}

```

```

# Store all parameters into an output variable

output <- list(p = p.t, mu = mu.t, sigma2 = sigma2.t, lambda = lambda.t)
attributes(output)$iter <- t

if (t == maxit) {
  warning('Reached maximum number of iterations')
  return(output)
} else if (sum(abs(params.prev - c(p.t, mu.t, sigma2.t, lambda.t))) < eps) {
  # Convergence check: sum|theta^(t) - theta^(t+1)| < epsilon (over i)
  return(output)
}
}
}

# Apply the EM algorithm to dataex5 to find the MLEs
theta <- em(dataex5, p.t = 0.1, mu.t = 1, sigma2.t = 0.25, lambda.t = 2)
theta

```

```

## $p
## [1] 0.4795454
##
## $mu
## [1] 2.013274
##
## $sigma2
## [1] 0.8637291
##
## $lambda
## [1] 1.033006
##
## attr("iter")
## [1] 519

```

- We will now draw the histogram of the data with estimated density superimposed:

```

load('dataex5.Rdata')

# Get params from MLE theta
p = theta[[1]]; mu = theta[[2]]; sigma2 = theta[[3]]; lambda = theta[[4]]

# Histogram of data with estimated density superimposed
hist(dataex5, breaks = 35, main="Random Sample from Mixture Distribution",
     xlab = "y",
     cex.main = 1.5,
     col= "lightblue",
     ylim= c(0,0.20),
     freq = FALSE)
# Superimposing mixed density distribution
curve(p*dlnorm(x, meanlog = mu, sdlog= sqrt(sigma2)) + (1-p)*dexp(x, lambda),
     add= TRUE, col= "red",lwd=1.5)
legend("topright", c("Histogram Data", "Mixed Distribution density"),
     fill=c("lightblue", "red"))

```

## Random Sample from Mixture Distribution

