# Incomplete Data Analysis Assignment 2

Josephine Li s2346729

# **Question 1**

#### Sub Question 1-a

First, we try to obtain the density function of Z, which is  $f_Z$ . First, calculate the cumulative distribution function of Z, which is  $F_Z$ . The cdf of Z with  $z \ge 1$  and  $\lambda, \mu > 0$  is:

$$\begin{split} F_Z(z) &= P(Z \leq z) \\ &= P(\min\{X,Y\} \leq z) \\ &= 1 - P(\min\{X,Y\} > z) \\ &= 1 - P(X > z, Y > z) \\ &= 1 - P(X > z) \times p(Y > z) \\ &= 1 - (1 - P(X \leq z)) \times (1 - P(Yleqz)) \\ &= 1 - (1 - F_X(z)) \times (1 - F_Y(z)) \\ &= 1 - z^{-(\lambda + \mu)} \end{split}$$

And the density function of Z with  $z \ge 1$  and  $\lambda, \mu > 0$  is:

$$f_Z(z) = \frac{d}{dz} F_Z(z) = \frac{\lambda + \mu}{z^{\lambda + \mu + 1}}$$

Then, try to obtain the frequency function of  $\delta$ , which is  $f_{\delta}$ . From 2 cumulative density function, we can calculate the density functions of X and Y with  $x, y \ge 1$  and  $\lambda, \mu > 0$ , which is:

$$f_X(x;\lambda) = \frac{d}{dx} F_X(x) = \frac{\lambda}{x^{\lambda+1}}$$
$$f_Y(y;\mu) = \frac{d}{dy} F_Y(y) = \frac{\mu}{y^{\mu+1}}$$

As 2 distributions are independent, he joint density function of 2 distribution X, Y is:

$$f_{XY}(x, y; \lambda, \mu) = f_X(x; \lambda) \times f_Y(y; \mu) = \frac{\lambda \times \mu}{x^{\lambda+1} \times y^{\mu+1}}$$

To calculate the frequency when  $\delta = 1$ , we need to calculate P(X < Y), which is:

$$P(X < Y) = \int_{1}^{\infty} \int_{x}^{\infty} \frac{\lambda \times \mu}{x^{\lambda + 1} \times y^{\mu + 1}} dy dx = \frac{\lambda}{\lambda + \mu}$$

And the frequency when  $\delta = 0$  is:

$$P(X \ge Y) = 1 - P(X < Y) = \frac{\mu}{\lambda + \mu}$$

Therefore, the frequency function of  $\delta$  is:

$$f_{\delta} = \begin{cases} \frac{\lambda}{\lambda + \mu}, & \delta = 1\\ \frac{\mu}{\lambda + \mu}, & \delta = 0 \end{cases}$$

The distribution of Z is Pareto-distribution with parameter  $\lambda + \mu$ , and the distribution of  $\delta$  is Bernoulli distribution with parameter  $\frac{\lambda}{\lambda + \mu}$ .

#### Sub Question 1-b

For samples  $Z_1, Z_2, \ldots, Z_n$  from  $f_Z(z; \theta)$  with  $\theta = \lambda + \mu$ , the joint density of observation  $\mathbb{Z} = (z_1, z_2, \ldots, z_n)$  is:

$$f_Z(\mathbf{z};\theta) = \prod_{i=1}^n f_Z(z_i;\theta) = \theta^n \prod_{i=1}^n z_i^{-(\theta+1)} = L(\theta;\mathbf{z})$$

 $L(\theta; z)$  is the likelihood function, and the log likelihood is:

$$\log L(\theta; z) = \log \theta^n \prod_{i=1}^n z_i^{-(\theta+1)}$$

$$= \log \theta^n + \log \prod_{i=1}^n z_i^{-(\theta+1)}$$

$$= n \log \theta - (\theta+1) \sum_{i=1}^n \log z_i$$

The maximum likelihood estimator of  $\theta$  is:

$$\hat{\theta}_{MLE} = \arg\max_{\theta \in \Theta} L(\theta; z)$$

To calculate the value of  $\hat{\theta}_{MLE}$ , we need to calculate the first derivatives for log likelihood function and let it equal to zero, which is:

$$\frac{d}{d\theta}\log L(\theta; \mathbf{z}) = \frac{n}{\theta} - \sum_{i=1}^{n} \log z_i = 0$$

We get the result  $\hat{\theta}_{MLE} = \frac{n}{\sum_{i=1}^{n} \log z_i}$ . To ensure that we have obtained a maximum, we need to confirm the derivative of the score function is negative. The derivative of the log likelihood function is:

$$\frac{d^2}{d\theta d\theta^T} \log L(\theta; \mathbf{z}) = -\frac{n}{\theta^2}$$

As  $z_i \geq 1$ , the second derivative is negative. Hence, the value of  $\hat{\theta}_{MLE}$  is:

$$\hat{\theta}_{MLE} = \frac{n}{\sum_{i=1}^{n} \log z_i}$$

We can do the same steps for deriving maximum likelihood estimator for p.

For samples  $\delta_1, \delta_2, \dots, \delta_n$  from  $f_{\delta}(d; p)$  with  $p = \frac{\lambda}{\lambda + \mu}$ , the probability mass function for  $\delta = (d_1, d_2, \dots, d_n)$  is:

$$f_{\delta}(\mathbf{d}; p) = p^d (1 - p)^{1 - d}, d \in \{0, 1\}$$

The likelihood function for p is:

$$L(p; \mathbf{d}) = \prod_{i=1}^{n} p^{d_i} (1-p)^{1-d_i} = p^{\sum_{i=1}^{n} d_i} \times (1-p)^{n-\sum_{i=1}^{n} d_i}$$

$$\log L(p; \mathbf{d}) = \log(p^{\sum_{i=1}^{n} d_i} \times (1-p)^{n-\sum_{i=1}^{n} d_i}) = \sum_{i=1}^{n} d_i \log p + (n - \sum_{i=1}^{n} d_i) \log(1-p)$$

Calculating the first derivatives for log likelihood function and let it equal to zero, which is:

$$\frac{d}{d\theta} \log L(p; d) = \frac{1}{p} \sum_{i=1}^{n} d_i - \frac{1}{1-p} (n - \sum_{i=1}^{n} d_i) = 0$$

We get the result  $\hat{p}_{MLE} = \frac{\sum_{i=1}^{n} d_i}{n} = \overline{\delta}$ . To ensure that we have obtained a maximum, we need to confirm the derivative of the score function is negative. The derivative of the log likelihood function is:

$$\frac{d^2}{dpdp^T}\log L(p; \mathbf{d}) = -\frac{1}{p^2} \sum_{i=1}^n d_i - \frac{1}{(1-p)^2} (n - \sum_{i=1}^n d_i)$$

As  $d_i \in \{0, 1\}$ , n is counting number of  $d_i$ , therefore,  $n \ge \sum_{i=1}^n$ . And  $p^2$ ,  $(1-p)^2 > 0$ , we can conclude that the second derivative is negative. Hence, the value of  $\hat{p}_{MLE}$  is:

$$\hat{p}_{MLE} = \frac{\sum_{i=1}^{n} d_i}{n} = \overline{\delta}$$

#### Sub Question 1-c

By the asymptotic normality of the MLE, we know that:

- for large sample sizes, the MLE follows an approximate normal distribution;
- the mean of the distribution is equal to the true value of the parameter being estimated;
- the standard deviation of the distribution is approximately equal to the inverse of the Fisher information, which can be estimated using the observed information.

For  $\theta$ , the expected Fisher information matrix is defined as:

$$I(\theta) = -E\left[\frac{d^2}{d\theta d\theta^2} \log L(\theta; z)\right] = \frac{n}{\theta^2}$$

The observed second derivatives evaluated at the MLE is:

$$J(\hat{\theta}_{MLE}) = \frac{n}{\hat{\theta}^2} = \frac{(\sum_{i=1}^n \log z_i)^2}{n}$$

As we known from the asymptotic normality of the MLE, we know that:

$$\hat{\theta} \sim N(\theta, J(\hat{\theta}_{MLE})^{-1})$$
$$\hat{\theta} \sim N(\hat{\theta}, \frac{\theta^2}{n})$$

Therefore, a 95% confidence ( $\alpha = 0.95$ ) interval for  $\theta$  is given by:

$$[\hat{\theta} - Z_{0+(1-0.95)/2} \times \sqrt{\frac{\theta^2}{n}}, \hat{\theta} + Z_{1-(1-0.95)/2} \times \sqrt{\frac{\theta^2}{n}}]$$

Which can be simplified as:

$$[\hat{\theta} - 1.96 \frac{\theta}{\sqrt{n}}, \hat{\theta} + 1.96 \frac{\theta}{\sqrt{n}}]$$

For p, the expected Fisher information matrix is defined as:

$$I(p) = -E\left[\frac{d^2}{dpdp^2}\log L(p; \mathbb{d})\right] = \frac{1}{p^2} \sum_{i=1}^n d_i + \frac{1}{(1-p)^2} (n - \sum_{i=1}^n d_i)$$

The observed second derivatives evaluated at the MLE is:

$$J(\hat{p}_{MLE}) = \frac{1}{\hat{p}^2} \sum_{i=1}^{n} d_i + \frac{1}{(1-\hat{p})^2} (n - \sum_{i=1}^{n} d_i)$$
$$= \frac{n}{\delta} + \frac{n}{1-\delta}$$
$$= \frac{n}{\delta(1-\delta)}$$

As we known from the asymptotic normality of the MLE, we know that:

$$\hat{p} \sim N(p, J(\hat{p}_{MLE})^{-1})$$

$$\hat{p} \sim N(\hat{p}, \frac{\overline{\delta}(1 - \overline{\delta})}{n})$$

Therefore, a 95% confidence ( $\alpha = 0.95$ ) interval for  $\theta$  is given by:

$$[\hat{p} - Z_{0+(1-0.95)/2} \times \sqrt{\frac{\overline{\delta}(1-\overline{\delta})}{n}}, \hat{p} + Z_{1-(1-0.95)/2} \times \sqrt{\frac{\overline{\delta}(1-\overline{\delta})}{n}}]$$

Which can be simplified as:

$$[\overline{\delta} - 1.96\sqrt{\frac{\overline{\delta}(1-\overline{\delta})}{n}}, \overline{\delta} + 1.96\sqrt{\frac{\overline{\delta}(1-\overline{\delta})}{n}}]$$

# **Question 2**

## Sub Question 2-a

The likelihood function of the uncensored data is given by:

$$L(\mu, \sigma^2 | y) = \prod_{i=1}^n f_Y(y_i; \mu, \sigma^2)$$

where  $f_Y(y_i; \mu, \sigma^2)$  is the probability density function of Y.

For the censored observations, we have that  $x_i = D$  if  $y_i < D$ , and  $x_i = y_i$  if  $y_i \ge D$ . The probability of observing  $x_i = D$  is equal to the probability of  $y_i < D$ , which is given by  $F_Y(D; \mu, \sigma^2)$ . Therefore, the likelihood function for the censored data is:

$$L(\mu, \sigma^2 | x, r) = \prod_{i=1}^n [f_X(x_i; \mu, \sigma^2)]^{r_i} \times [F_Y(D; \mu, \sigma^2)]^{1-r_i}$$

where  $f_X(x_i; \mu, \sigma^2)$  is the probability density function of X, which is equal to:

$$f_X(x_i; \mu, \sigma^2) = \begin{cases} f_Y(x_i; \mu, \sigma^2) & \text{if } x_i \ge D \\ F_Y(D; \mu, \sigma^2) & \text{if } x_i < D \end{cases}$$

Taking the logarithm of this expression yields the log-likelihood function:

$$\log L(\mu, \sigma^{2}; x, r) = \sum_{i=1}^{n} \{ r_{i} \log f_{X}(x_{i}; \mu, \sigma^{2}) + (1 - r_{i}) \log F_{Y}(D; \mu, \sigma^{2}) \}$$

$$= \sum_{i=1}^{n} \{ r_{i} \log \phi(x_{i}; \mu, \sigma^{2}) + (1 - r_{i}) \log \Phi(x_{i}; \mu, \sigma^{2}) \}$$

#### **Sub Question 2-b**

The log likelihood for  $\mu$  and  $\sigma^2$  is given by:

$$\log L(\mu, \sigma^{2}; x, r) = \sum_{i=1}^{n} \{ r_{i} \log \phi(x_{i}; \mu, \sigma^{2}) + (1 - r_{i}) \log \Phi(x_{i}; \mu, \sigma^{2}) \}$$

$$= \sum_{i=1}^{n} \{ r_{i} \log(\frac{1}{\sigma\sqrt{2\pi}} e^{-\frac{(x_{i} - \mu)^{2}}{2\sigma^{2}}}) + (1 - r_{i}) \log(\frac{1}{\sigma\sqrt{2\pi}} \int_{-\infty}^{x_{i}} e^{-\frac{(x_{i} - \mu)^{2}}{2\sigma^{2}}}) \}$$

As we already known  $\sigma 2 = 1.5^2$ , the MLE value for  $\mu$  is given by:

$$\frac{d}{d\mu}\log L(\mu; x, r) = 0$$

Read data from dataex2.Rdata file.

```
load('dataex2.Rdata')
# library(maxLik)
require(maxLik)
```

```
## Loading required package: maxLik
```

```
## Loading required package: miscTools
```

```
##
## Please cite the 'maxLik' package as:
## Henningsen, Arne and Toomet, Ott (2011). maxLik: A package for maximum likelihood
estimation in R. Computational Statistics 26(3), 443-458. DOI 10.1007/s00180-010-0217
-1.
##
## If you have questions, suggestions, or comments regarding the 'maxLik' package, pl
ease use a forum or 'tracker' at maxLik's R-Forge site:
## https://r-forge.r-project.org/projects/maxlik/
```

```
log_like <- function(mu,data,sd=1.5){
    x <- data[1]
    r <- data[2]
    sum(r*log(dnorm(as.matrix(x),mean = mu,sd))
        + (1-r)*log(pnorm(as.matrix(x),mean = mu,sd )))
}
mle <- maxLik(logLik = log_like,data = dataex2, start = c(mu=5))
summary(mle)</pre>
```

The maximum likelihood estimator for  $\mu$  is 5.5328.

## **Question 3**

#### Sub Question 3-a

The missing data mechanism can be disregarded for likelihood inference, given that it is MAR (Missing At Random) and only associated with the observed value of  $Y_1$ , while  $\psi$  is different from  $\theta$ . The missing mechanism can be represented as:

$$L(\theta, \psi | \mathbf{y}_{obs}, \mathbf{r}) = f(\mathbf{r} | \mathbf{y}_{obs}, \psi) L(\theta | \mathbf{y}_{obs})$$

Therefore, it is ignoreable for likelihood estimation.

## Sub Question 3-b

The missing data mechanism cannot be ignored for likelihood inference, as it is MNAR (Missing Not At Random) and associated with the missing value of  $Y_2$ , while  $\psi$  is different from  $\theta$ . The missing mechanism can be represented as:

$$L(\theta, \psi | \mathbf{y}_{mis}, \mathbf{r}) = f(\mathbf{r} | \mathbf{y}_{obs}, \psi) L(\theta | \mathbf{y}_{obs})$$

Therefore, it is non-ignoreable for likelihood estimation.

## Sub Question 3-c

The missing data mechanism cannot be disregarded for likelihood inference, as it is MAR (Missing At Random) and associated only with the observed value of  $Y_1$ , but not distinct from  $\theta$ . The mechanism can be represented as:

$$L(\theta, \psi | \mathbf{y}_{obs}, \mathbf{r}) = f(\mathbf{r} | \mathbf{y}_{obs}, \psi, \mu_1) L(\theta | \mathbf{y}_{obs})$$

Therefore, it is ignoreable for likelihood estimation.

# **Question 4**

We calculate the log likelihood for  $\beta$  of the complete data first.

As  $Y_i \sim Bernoulli\{p_i(\beta)\}\$ , the density function of Y is:

$$f(Y_i; \beta) = p_i(\beta)^{Y_i} (1 - p_i(\beta))^{1 - Y_i}$$

with  $p_i(\beta) = (exp(\beta_0 + \beta_1)) \div (1 + exp(\beta_0 + \beta_1))$ , the density function is:

$$f(Y_i; \beta) = \left[\frac{exp(\beta_0 + x_i\beta_1)}{1 + exp(\beta_0 + x_i\beta_1)}\right]^{Y_i} \times \left[\frac{1}{1 + exp(\beta_0 + x_i\beta_1)}\right]^{1 - Y_i}$$

the complete likelihood function is:

$$L(\beta; Y_i) = \prod_{i=1}^n f(Y_i; \beta)$$

$$= \prod_{i=1}^n \left[ \frac{exp(\beta_0 + x_i\beta_1)}{1 + exp(\beta_0 + x_i\beta_1)} \right]^{Y_i} \times \left[ \frac{1}{1 + exp(\beta_0 + x_i\beta_1)} \right]^{1-Y_i}$$

the complete log likelihood function is:

$$\begin{split} \log L(\beta; Y_i) &= \prod_{i=1}^n \log f(Y_i; \beta) \\ &= \prod_{i=1}^n \log \{ [\frac{exp(\beta_0 + x_i\beta_1)}{1 + exp(\beta_0 + x_i\beta_1)}]^{Y_i} \times [\frac{1}{1 + exp(\beta_0 + x_i\beta_1)}]^{1 - Y_i} \} \\ &= \sum_{i=1}^n Y_i(\beta_0 + x_i\beta_1) - \sum_{i=1}^n \log (1 + exp(\beta_0 + x_i\beta_1)) \end{split}$$

Then, applying the EM algorithm to estimate the values of  $\beta$ .

For **E-step** at iteration t+1, we need to calculate the expectation, with respect to what is missing,  $Y_i$ , of the above complete data likelihood, given what is observed  $x_i$  and the current estimate of  $\beta$ , that is:

$$\begin{split} Q(\beta|\beta^{t}) &= E_{Y}[\log L(\beta|Y,x)|Y_{obs},x,\beta^{t}] \\ &= E_{Y_{m+1}...m}[\log L(\beta|Y,x)|Y_{1..m},x_{1..m},\beta^{t}] \\ &= E[\sum_{i=1}^{m} Y_{i}(\beta_{0} + x_{i}\beta_{1}) - \sum_{i=1}^{m} \log(1 + exp(\beta_{0} + x_{i}\beta_{1})) \\ &+ \sum_{i=m+1}^{n} Y_{i}(\beta_{0} + x_{i}\beta_{1}) - \sum_{i=m+1}^{n} \log(1 + exp(\beta_{0} + x_{i}\beta_{1}))] \\ &= \sum_{i=1}^{m} Y_{i}(\beta_{0} + x_{i}\beta_{1}) - \sum_{i=1}^{n} \log(1 + exp(\beta_{0} + x_{i}\beta_{1})) \\ &+ \sum_{i=m+1}^{n} \frac{exp(\beta_{0}^{t} + x_{i}\beta_{1}^{t})}{1 + exp(\beta_{0}^{t} + x_{i}\beta_{1}^{t})} (\beta_{0} + x_{i}\beta_{1}) \end{split}$$

where the first m value of Y is observed and m + 1..n value of Y is missing.

For **M-step**, we obtain  $\beta^{t+1}$ , the value of  $\beta$  that maximize  $Q(\beta|\beta^t)$ .

To calculate  $\beta^{t+1}$ , we need to do first partial derivatives for  $\beta_0, \beta_1$  and let them equal to 0, which is:

$$\begin{split} \frac{\partial}{\partial \beta_0} Q &= \frac{\partial}{\partial \beta_0} \big[ \sum_{i=1}^m Y_i (\beta_0 + x_i \beta_1) - \sum_{i=1}^n \log(1 + \exp(\beta_0 + x_i \beta_1)) \\ &+ \sum_{i=m+1}^n \frac{\exp(\beta_0^t + x_i \beta_1^t)}{1 + \exp(\beta_0^t + x_i \beta_1^t)} (\beta_0 + x_i \beta_1) \big] \\ &= \sum_{i=1}^m Y_i - \sum_{i=1}^n \frac{\exp(\beta_0 + x_i \beta_1)}{1 + \exp(\beta_0 + x_i \beta_1)} + \sum_{i=m+1}^n \frac{\exp(\beta_0^t + x_i \beta_1^t)}{1 + \exp(\beta_0^t + x_i \beta_1^t)} = 0 \\ \frac{\partial}{\partial \beta_1} Q &= \frac{\partial}{\partial \beta_1} \big[ \sum_{i=1}^m Y_i (\beta_0 + x_i \beta_1) - \sum_{i=1}^n \log(1 + \exp(\beta_0 + x_i \beta_1)) \\ &+ \sum_{i=m+1}^n \frac{\exp(\beta_0^t + x_i \beta_1^t)}{1 + \exp(\beta_0^t + x_i \beta_1^t)} (\beta_0 + x_i \beta_1) \big] \\ &= \sum_{i=1}^m Y_i x_i - \sum_{i=1}^n \frac{\exp(\beta_0 + x_i \beta_1)}{1 + \exp(\beta_0 + x_i \beta_1)} x_i + \sum_{i=m+1}^n \frac{\exp(\beta_0^t + x_i \beta_1^t)}{1 + \exp(\beta_0^t + x_i \beta_1^t)} x_i = 0 \end{split}$$

Besides, we need to make sure that the second derivatives are negative.

$$\frac{\partial^2}{\partial \beta_0^2} Q = -\sum_{i=1}^n \frac{\exp(\beta_0 + x_i \beta_1)}{(1 + \exp(\beta_0 + x_i \beta_1))^2} - \sum_{i=m+1}^n \frac{\exp(\beta_0^t + x_i \beta_1^t) x_i^2}{(1 + \exp(\beta_0^t + x_i \beta_1^t))^2}$$
$$\frac{\partial^2}{\partial \beta_1^2} Q = -\sum_{i=1}^n \frac{x_i^2 \exp(\beta_0 + x_i \beta_1)}{(1 + \exp(\beta_0 + x_i \beta_1))^2} - \sum_{i=m+1}^n \frac{\exp(\beta_0^t + x_i \beta_1^t) x_i^2}{(1 + \exp(\beta_0^t + x_i \beta_1^t))^2}$$

Therefore, we can obtain the values of  $\beta_0^{t+1}, \beta_1^{t+1}$  from the first derivatives when they equal to 0.

Then, repeating M-step and E step until some convergence criterion is met. Here we set criterion like:

$$|\beta_0^{t+1} - \beta_0^t| + |\beta_1^{t+1} - \beta_1^t| \le 0.0001$$

Then, we try to encode EM algorithm and calculate the values of  $\beta$  .

```
# workshop4
load('dataex4.Rdata')
EMex4 <- function(beta ini, data EM, eps){</pre>
diff <- 1
beta <- beta ini
i <- 0
while(diff > eps){
beta.old <- beta
mle_beta <- maxLik(logLik = log_like_ex4, data =</pre>
                      data EM, beta t = beta.old,start = beta ini)
beta <- mle beta$estimate
diff <- sum(abs(beta-beta.old))</pre>
i <- i+1
}
return(beta)
beta= EMex4(beta_ini = c(0, 0), data_EM = dataex4,0.0001)
```

```
## [1] 0.9755477 -2.4802557
```

The result for  $\beta_0$  is 0.9755, for  $\beta_1$  is -2.4803.

## Question 5

#### Sub Question 5-a

We calculate the log likelihood for  $\theta$  of the complete data first. According to the question, CDF of  $F_Y$  is:

$$F(y; \lambda, \mu, p) = p \times F_X(y; \lambda) + (1 - p) \times F_Y(y; \mu)$$
  
=  $p \times (1 - y^{-\lambda}) + (1 - p) \times (1 - y^{-\mu})$   
=  $1 - p \times y^{-\lambda} - y^{-\mu} + p \times y^{-\mu}$ 

with  $y \ge 1$  and  $\lambda, \mu > 0$ .

Do the first derivatives to calculate PDF of Y, which is:

$$f_Y(y; \lambda, \mu, p) = \frac{\partial}{\partial y} F(y; \lambda, \mu, p)$$

$$= p \times y^{-(\lambda+1)} + y^{-(\mu+1)} - p \times y^{-(\mu+1)}$$

$$= p \times \lambda \times y^{-(\lambda+1)} + (1-p) \times \mu \times y^{-(\mu+1)}$$

Define Z to be the indicator of Y, the rule is:

$$Z_i = \begin{cases} 1, & \text{if } Y_i \text{ is from } F_X \\ 0, & \text{if } Y_i \text{ is from } F_Y \end{cases} \quad f_Z = \begin{cases} p, Z_i = 1 \\ 1 - p, Z_i = 0 \end{cases}$$

The PDF of Y and Z can be written as:

$$f(y, z; \lambda, \mu, p) = [p \times y^{-(\lambda+1)}]^{z_i} + [y - (\mu + 1) - p \times y - (\mu + 1)]^{1-z_i}$$

The likelihood function can be written as:

$$L(\theta; \mathbf{y}, \mathbf{z}) = \prod_{i=1}^{n} f(y, z; \lambda, \mu, p)$$
$$= \prod_{i=1}^{n} [p \times \lambda \times y^{-(\lambda+1)}]^{z_i} \times [(1-p) \times \mu \times y^{-(\mu+1)}]^{(1-z_i)}$$

The log likelihood function is:

$$\log L = \sum_{i=1}^{n} z_i \log[p \times \lambda \times y^{-(\lambda+1)}] + \sum_{i=1}^{n} (1 - z_i) \log[(1 - p) \times \mu \times y^{-(\mu+1)}]$$

Then, applying the EM algorithm to estimate the values of  $\theta$ .

For **E-step** at iteration t+1, we need to calculate the expectation, with respect to what is from  $F_X$  or  $F_Y$  of the above complete data likelihood, given what is observed  $y_i$  and the current estimate of  $\theta$ , that is:

$$\begin{split} Q(\theta|\theta^t) &= E_z[\log L(\theta|\mathbf{y},z)|\mathbf{y},\theta^t] \\ &= \sum_{i=1}^n E[z_i|\mathbf{y},\theta^t] \times \log(p \times \lambda \times y^{-(\lambda+1)}) \\ &+ \sum_{i=1}^n (1 - E[z_i|\mathbf{y},\theta^t]) \times \log((1-p) \times \mu \times y^{-(\mu+1)}) \end{split}$$

with expectation of Z, which is  $E[z_i|\mathbf{y},\theta^t]$ , can be calculated as:

$$\begin{split} E[z_i|\mathbf{y},\theta^t] &= 1 \times p_Z(Z=1) + 0 \times (1-p_Z(Z=1)) \\ &= \frac{p^t \times \lambda^t \times y^{-(\lambda^t+1)}}{p^t \times \lambda^t \times y^{-(\lambda^t+1)} + (1-p^t) \times \mu^t \times y^{-(\mu^t+1)}} \end{split}$$

Using  $Q(\theta|\theta^t)$  and  $E[z_i|y,\theta^t]$ , we can complete our calculation in E step.

For **M-step**, we obtain  $\theta^{t+1}$ , the value of  $\theta$  that maximise  $Q(\theta|\theta^t)$ .

To calculate  $\theta^{t+1}$ , we need to do first partial derivatives for  $\lambda, \mu, p$  and let them equal to 0, which is:

$$\frac{\partial}{\partial p} Q(\theta | \theta^t) = \frac{\sum_{i=1}^n E[z_i | \mathbf{y}, \theta^t]}{p} = 0$$

$$\frac{\partial}{\partial \lambda} Q(\theta | \theta^t) = \frac{\sum_{i=1}^n E[z_i | \mathbf{y}, \theta^t]}{\lambda} - \sum_{i=1}^n E[z_i | \mathbf{y}, \theta^t] \times log(y_i) = 0$$

$$\frac{\partial}{\partial \mu} Q(\theta | \theta^t) = \frac{\sum_{i=1}^n (1 - E[z_i | \mathbf{y}, \theta^t])}{\mu} - \sum_{i=1}^n (1 - E[z_i | \mathbf{y}, \theta^t]) \times log(y_i) = 0$$

Check the second derivatives:

$$\frac{\partial^2}{\partial \rho \partial \rho^T} Q(\theta | \theta^t) = -\frac{\sum_{i=1}^n E[z_i | \mathbf{y}, \theta^t]}{\rho^2}$$
$$\frac{\partial^2}{\partial \lambda \partial \lambda^T} Q(\theta | \theta^t) = -\frac{\sum_{i=1}^n E[z_i | \mathbf{y}, \theta^t]}{\lambda^2}$$
$$\frac{\partial^2}{\partial \mu \partial \mu^T} Q(\theta | \theta^t) = -\frac{\sum_{i=1}^n (1 - E[z_i | \mathbf{y}, \theta^t])}{\mu^2}$$

The second derivatives are all negative, therefore, we can obtain  $p^{t+1}$ ,  $\lambda^{t+1}$ ,  $\mu^{t+1}$ , which are:

$$p^{t+1} = \frac{\sum_{i=1}^{n} E[z_i | \mathbf{y}, \theta^t]}{n} = E[z_i | \mathbf{y}, \theta^t]$$
$$\lambda^{t+1} = \frac{\sum_{i=1}^{n} E[z_i | \mathbf{y}, \theta^t]}{\sum_{i=1}^{n} E[z_i | \mathbf{y}, \theta^t] \times \log(y_i)}$$
$$\mu^{t+1} = \frac{\sum_{i=1}^{n} (1 - E[z_i | \mathbf{y}, \theta^t]) \times \log(y_i)}{\sum_{i=1}^{n} (1 - E[z_i | \mathbf{y}, \theta^t]) \times \log(y_i)}$$

Then, repeating M-step and E step until some convergence criterion is met. Here we set criterion like:

$$\left|\lambda^{t+1} - \lambda^{t}\right| + \left|p^{t+1} - p^{t}\right| + \left|\mu^{t+1} - \mu^{t}\right| \le 0.0001$$

### Sub Question 5-b

In this part, try to encode EM algorithm and calculate the values of  $\theta$  .

```
rm(list = ls())
EMex5 <- function(ini theta,data,eps){</pre>
  diff <- 1
  y <- data
  theta <- ini theta
  p <- theta[1]; lambda <- theta[2]; mu <- theta[3]</pre>
  n <- length(y)</pre>
  while(diff > eps){
    theta.old <- theta
    #E-step
    p1 <- p*lambda*y**(-lambda-1)</pre>
    p2 < (1-p)*mu*y**(-mu-1)
    p_hat <- p1/(p1+p2)
    #M-step
    p <- sum(p hat)/n
    lambda <- sum(p_hat)/sum(p_hat*log(y))</pre>
    mu <- sum(1-p_hat)/sum((1-p_hat)*log(y))
    #theta(t+1)
    theta <- c(p, lambda, mu)
    diff <- sum(abs(theta-theta.old))</pre>
  }
  return(theta)
load("dataex5.Rdata")
theta <- EMex5(ini theta = c(0.3,0.3,0.4), data = dataex5, 0.0001)
p \leftarrow theta[1]
lambda <- theta[2]</pre>
mu<- theta[3]
#estimated p; estimated lambda; estimated mu
theta
```

The estimate values from EM algorithm is p = 0.7939,  $\lambda = 0.9763$ ,  $\mu = 0.6706$ .

Then, drawing the histogram of the data with the estimated density superimposed.

```
#function to calculate density function of pareto distribution
dpareto <- function (x, k){
  f < -k*x**(-k-1)
  return(f)
}
load("dataex5.Rdata")
y <- dataex5
hist(y, breaks = "FD", main = "Histogram with Estimated Density",
     xlab = "dataex5",
     ylab = "Density",
     cex.main = 1.5, cex.lab = 1.5, cex.axis = 1.4,
     freq = F, ylim = c(0,1.5), xlim = c(0,15))
#add the estimated density curve
x <- y
curve(p*dpareto(x,lambda)+(1-p)*dpareto(x,mu),
      add = TRUE, lwd = 2, col = "red")
```

# **Histogram with Estimated Density**

