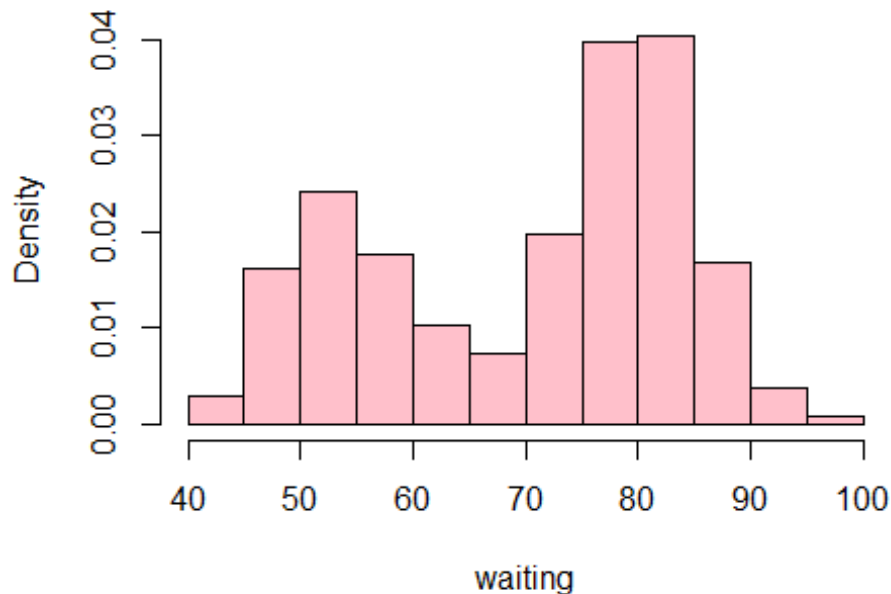


Problem 3: Analysis of faithful datasets.

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Consider the faithful datasets:

```
attach(faithful)
waiting = sort(waiting)
hist(faithful$waiting,xlab = 'waiting',probability = T,col='pink',main='')
```



Fit following three models using MLE method and calculate **Akaike information criterion** (aka., AIC) for each fitted model. Based on AIC decides which model is the best model?

Based on the best model calculate the following probability

$$\mathbb{P}(60 < \text{waiting} < 70)$$

(i) **Model 1:**

$$f(x) = p * \text{Gamma}(x|\alpha, \sigma_1) + (1 - p)N(x|\mu, \sigma_2^2), \quad 0 < p < 1$$

Method 1

```
NegLogLike1 <- function(theta,data){
  alpha1 = exp(theta[1])
  beta1 = exp(theta[2])
  mu2 = theta[3]
```

```

sigma2 = exp(theta[4])
p = exp(theta[5])/(1+exp(theta[5]))
n = length(data)
l=0
for(i in 1:n){
  l = l + log(p*dgamma(data[i],shape = alpha1,scale= beta1)
             +(1-p)*dnorm(data[i], mean=mu2, sd= sigma2))
}
return(-l)
}
#Length(waiting[waiting<67])/ Length(waiting)
theta1_initial=c(log(89.05),log(0.61),80.21,log(5.70),-0.58)

#NegLogLikeMix(theta_initial,waiting)

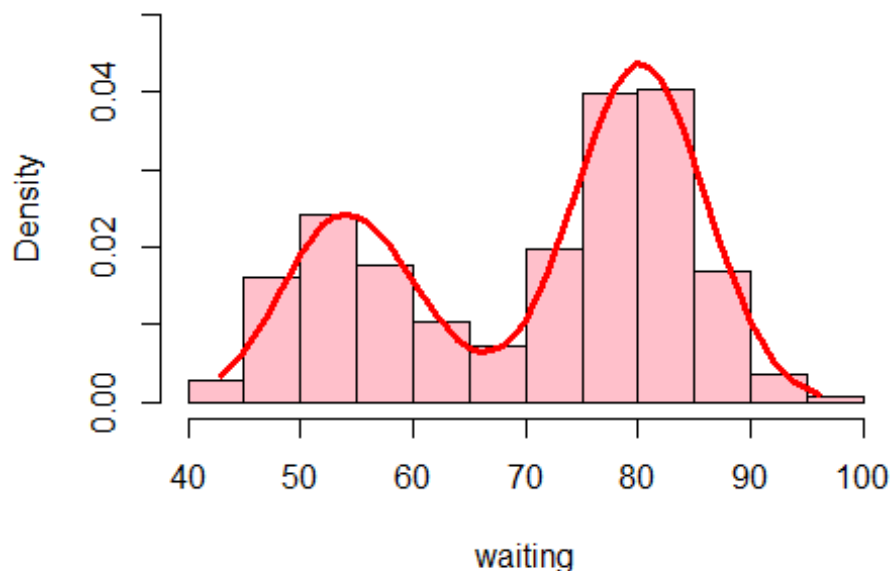
fit1 = optim(theta1_initial
             ,NegLogLike1
             ,data=waiting)

theta1_hat = fit1$par
alpha11_hat = exp(theta1_hat[1])
beta11_hat = exp(theta1_hat[2])
mu12_hat = theta1_hat[3]
sigma12_hat = exp(theta1_hat[4])
p1_hat = exp(theta1_hat[5])/(1+exp(theta1_hat[5]))

d_mle1 = p1_hat*dgamma(waiting,shape = alpha11_hat, scale = beta11_hat)+(1-
p1_hat)*dnorm(waiting,mean=mu12_hat,sd=sigma12_hat)

hist(faithful$waiting,xlab = 'waiting',
     probability = T,col='pink',main='',ylim=c(0.,0.05))
lines(waiting,d_mle1,lwd=3,col='red')

```



(ii) **Model 2:**

$$f(x) = p * \text{Gamma}(x|\alpha_1, \sigma_1) + (1 - p)\text{Gamma}(x|\alpha_2, \sigma_2), \quad 0 < p < 1$$

Method 2

```
NegLogLike2 <- function(theta,data){
  alpha1 = exp(theta[1])
  beta1 = exp(theta[2])
  alpha2 = exp(theta[3])
  beta2 = exp(theta[4])
  p = exp(theta[5])/(1+exp(theta[5]))
  n = length(data)
  l=0
  for(i in 1:n){
    l = l + log(p*dgamma(data[i],shape = alpha1,scale = beta1)
               +(1-p)*dgamma(data[i],shape=alpha2,scale =beta2))
  }
  return(-l)
}
#length(waiting[waiting<65])/ length(waiting)

#theta2_initial=c(log(78),log(1.5),log(104),log(1.3),-0.62) #wrong initial
values
theta2_initial=c(log(78),log(0.7),log(104),log(0.8),-0.62)

#theta_initial=c(log(89.05),log(0.61),log(198.12),log(0.4), -0.58) #doesn't
```

```

work interchange parameters and then works
#theta_initial = c(78,1.5,104,1.3,.35)
#NegLogLikeMix(theta_initial,waiting)

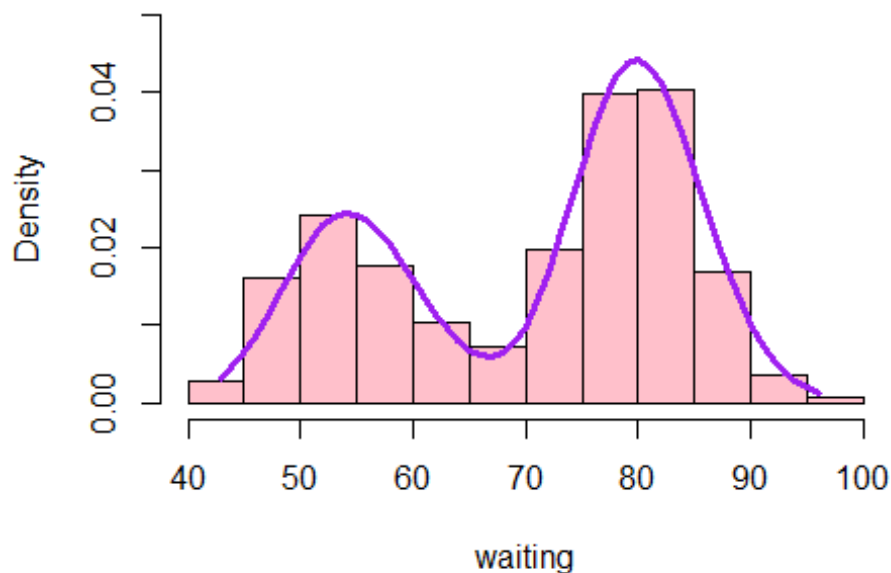
fit2 = optim(theta2_initial
            ,NegLogLike2
            ,data=waiting)

theta2_hat = fit2$par
alpha21_hat = exp(theta2_hat[1])
beta21_hat = exp(theta2_hat[2])
alpha22_hat = exp(theta2_hat[3])
beta22_hat = exp(theta2_hat[4])
p2_hat = exp(theta2_hat[5])/(1+exp(theta2_hat[5]))

d_mle2 = p2_hat*dgamma(waiting,shape = alpha21_hat,scale = beta21_hat) +(1-
p2_hat)*dgamma(waiting,shape=alpha22_hat,scale=beta22_hat)

hist(faithful$waiting,xlab = 'waiting',
     probability = T,col='pink',main='',ylim=c(0.,0.05))
lines(waiting,d_mle2,lwd=3,col='purple')

```



(iii) **Model 3:**

$$f(x) = p * \logNormal(x|\mu_1, \sigma_1^2) + (1 - p)\logNormal(x|\mu_1, \sigma_1^2), \quad 0 < p < 1$$

Method 3

```
NegLogLike3 <- function(theta,data){
  mu1 = exp(theta[1])
  sigma1 = exp(theta[2])
  mu2 = exp(theta[3])
  sigma2 = exp(theta[4])
  p = exp(theta[5])/(1+exp(theta[5]))
  n = length(data)
  l=0
  for(i in 1:n){
    l = l + log(p*dlnorm(data[i],mean = mu1, sd=sigma1)
               +(1-p)*dlnorm(data[i], mean=mu2, sd= sigma2))
  }
  return(-l)
}
#Length(waiting[waiting<65])/ Length(waiting)
theta3_initial=c(1.37,-2.16,1.48,-2.3,-0.62)

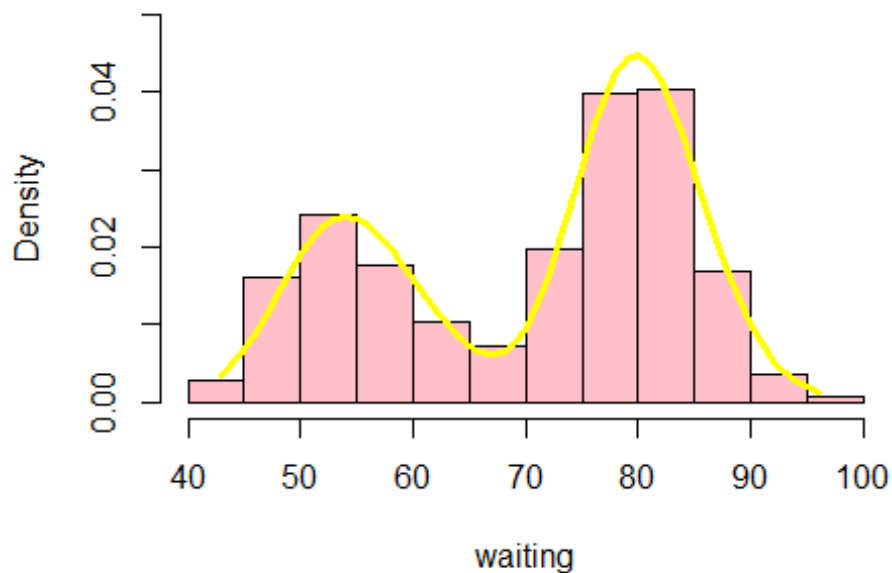
#NegLogLikeMix(theta_initial,waiting)

fit3 = optim(theta3_initial
             ,NegLogLike3
             ,data=waiting)

theta3_hat = fit3$par
mu31_hat = exp(theta3_hat[1])
sigma31_hat = exp(theta3_hat[2])
mu32_hat = exp(theta3_hat[3])
sigma32_hat = exp(theta3_hat[4])
p3_hat = exp(theta3_hat[5])/(1+exp(theta3_hat[5]))

d_mle3 = p3_hat*dlnorm(waiting,mean=mu31_hat, sd=sigma31_hat) +(1-
p3_hat)*dlnorm(waiting,mean=mu32_hat,sd=sigma32_hat)

hist(faithful$waiting,xlab = 'waiting',
     probability = T,col='pink',main='',ylim=c(0.,0.05))
lines(waiting,d_mle3,lwd=3,col='yellow')
```



```
# AIC

model = list(fit1,fit2,fit3)

frame=data.frame('Model'=c(0),'No_of_parameters'=c(50),
                  'Max_log_likelihood'=c(0))
for (i in 1:length(model)){

  #aic = 2*length(model[[i]]$par) + 2 *model[[i]]$value

frame=rbind(frame,c(paste('Model',as.character(i)),as.numeric(length(model[[i]]$par)),
                    -as.numeric(model[[i]]$value)))
}
frame = frame[-1,]
frame$AIC = 2*as.numeric(frame$No_of_parameters) -
2*as.numeric(frame$Max_log_likelihood)

abc = frame[frame$AIC == min(frame$AIC),]

frame[,c('Model', 'AIC')]
```

```

##      Model      AIC
## 2 Model 1 2076.180
## 3 Model 2 2076.156
## 4 Model 3 2075.420

cat(" The best model is : ",abc$Model )

## The best model is : Model 3

#Probability calc

dMix<-function(x,theta){
  mu1 = theta[1]
  sigma1 = theta[2]
  mu2 = theta[3]
  sigma2 = theta[4]
  p = theta[5]
  f = p*dlnorm(x,mean = mu1,sd=sigma1)+(1-p)*dlnorm(x,mean=mu2,sd=sigma2)
  return(f)
}

prob =
integrate(dMix,60,70,c(mu31_hat,sigma31_hat,mu32_hat,sigma32_hat,p3_hat))

print(paste("Probability (60 < waiting < 70) = ", prob$value))

## [1] "Probability (60 < waiting < 70) = 0.0906679402261568"

```