Computational Statistics

Lab 6

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Question 1

1.1

```
genfunc <- function(x) {
    (x^2 / exp(x)) - 2 * exp(-(9 * sin(x)) / (x^2 + x + 1))
}</pre>
```

1.2

```
crossover <- function(x,y) {
   (x + y) / 2
}</pre>
```

1.3

```
mutate <- function(x) {
    x^2 %% 30
}</pre>
```

1.4

```
genetic <- function(maxiter, mutprob) {
    ## a)
    ## plot(x = 0:30, y= genfunc(0:30), xlim = c(0,30), type ="l", xlab="", ylab="")

## b)
    X <- seq(0,30,by = 5)

## c)
    values <- genfunc(X)
    ## points(X, values, col = "red")

## d)
    bestvalue <- -Inf

for (i in 1:maxiter){
    ## i</pre>
```

```
parents <- sample(1:length(X), size = 2 )</pre>
    ## ii
    victim <- which.min(values)</pre>
    ## iii
    child <- crossover(X[parents[1]],X[parents[2]])</pre>
    if (mutprob > runif(1,0,1)) {
         child <- mutate(child)</pre>
    }
    ## iv
    X[victim] <- child</pre>
    values[victim] <- genfunc(child)</pre>
    ## values <- genfunc(X)</pre>
    ## v
    bestvalue <- max(bestvalue, max(values))</pre>
}
## points(x = X, y = values, col = "darkgreen")
list(opt=bestvalue, pop=X, vals=values)
```

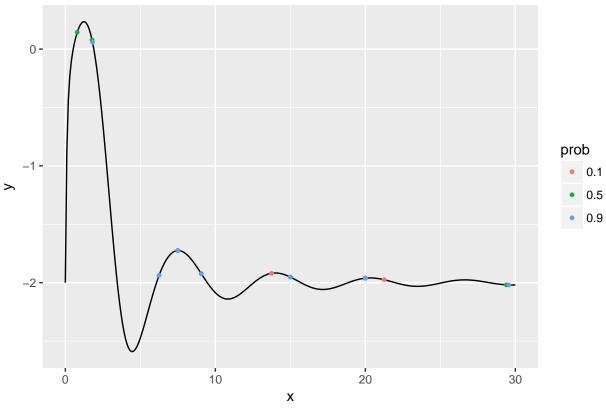
1.5

```
library(ggplot2)
func_{data} \leftarrow data.frame(x=seq(0, 30, by=0.1), y=genfunc(seq(0, 30, by=0.1)))
set.seed(123456)
r1 <- genetic(maxiter = 10, mutprob = 0.1)
r1$opt
## [1] -1.7
set.seed(123456)
r2 <- genetic(maxiter = 10, mutprob = 0.5)
r2$opt
## [1] 0.14
set.seed(123456)
r3 <- genetic(maxiter = 10, mutprob = 0.9)
r3$opt
## [1] 0.059
rd1 <- data.frame(x=r1$pop, y=r1$vals, prob="0.1")
rd2 <- data.frame(x=r2$pop, y=r2$vals, prob="0.5")
rd3 <- data.frame(x=r3$pop, y=r3$vals, prob="0.9")
```

```
plot_data <- rbind(rd1, rd2, rd3)

ggplot() +
    ggtitle("10 Iterations") +
    geom_line(data=func_data, aes(x=x, y=y)) +
    geom_point(data=plot_data, aes(x=x, y=y, col=prob), size=1) +
    theme(plot.title=element_text(hjust=0.5))</pre>
```

10 Iterations



```
set.seed(123456)
r1 <- genetic(maxiter = 100, mutprob = 0.1)
r1$opt

## [1] -1.7
set.seed(123456)
r2 <- genetic(maxiter = 100, mutprob = 0.5)
r2$opt

## [1] 0.23
set.seed(123456)
r3 <- genetic(maxiter = 100, mutprob = 0.9)
r3$opt

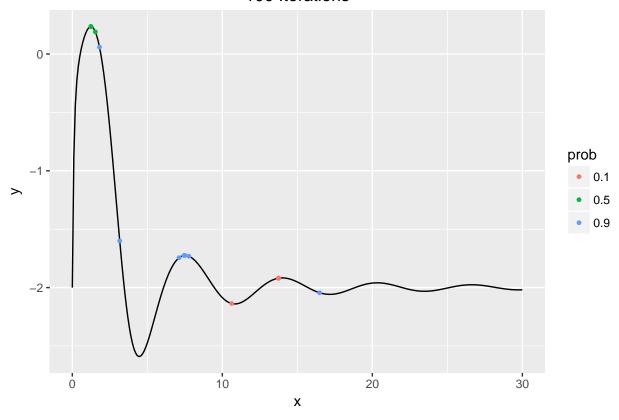
## [1] 0.059
rd1 <- data.frame(x=r1$pop, y=r1$vals, prob="0.1")
rd2 <- data.frame(x=r2$pop, y=r2$vals, prob="0.5")</pre>
```

```
rd3 <- data.frame(x=r3$pop, y=r3$vals, prob="0.9")

plot_data <- rbind(rd1, rd2, rd3)

ggplot() +
    ggtitle("100 Iterations") +
    geom_line(data=func_data, aes(x=x, y=y)) +
    geom_point(data=plot_data, aes(x=x, y=y, col=prob), size=1) +
    theme(plot.title=element_text(hjust=0.5))</pre>
```

100 Iterations

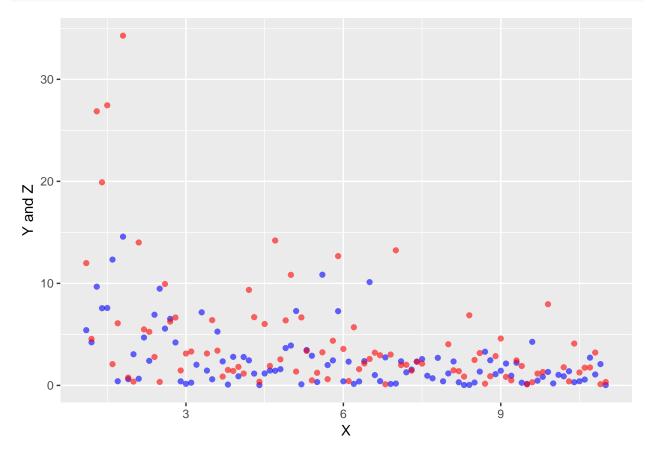


Question 2

2.1

```
physical <- read.csv("../data/physical1.csv")

ggplot(physical) +
    geom_point(aes(x=X, y=Y), col="blue", alpha = 0.6) +
    geom_point(aes(x=X, y=Z), col="red", alpha = 0.6) +
    labs(y = "Y and Z")</pre>
```



2.2

We know

$$Y_i \sim \exp(X_i/\lambda),$$

 $Z_i \sim \exp(X_i/(2\lambda)).$

To find λ using the EM-algorithm we want to find the likelihood which is

$$\begin{split} l(\lambda) &= P(\lambda|X,Y,Z) \\ &= \prod_{i=1}^{n} \frac{x_i}{\lambda} \exp\left(-\frac{x_i}{\lambda} y_i\right) \frac{x_i}{2\lambda} \exp\left(-\frac{x_i}{2\lambda} y_i\right) \\ &= \prod_{i=1}^{n} \frac{x_i}{2\lambda^2} \exp\left(-\frac{x_i}{\lambda} \left(y_i + \frac{z_i}{2}\right)\right). \end{split}$$

Since we know that multiplication with small numbers is bad for accuracy reasons in the computer we want the log-likelihood

$$L(\lambda) = \sum_{i=1}^{n} \log \left(\frac{x_i}{2\lambda^2} \exp\left(-\frac{x_i}{\lambda} \left(y_i + \frac{z_i}{2} \right) \right) \right)$$
$$= \sum_{i=1}^{n} \left(\log(x_i^2) - 2n \log(2\lambda) - \frac{x_i}{\lambda} \left(y_i + \frac{z_i}{2} \right) \right)$$
$$= 2 \sum_{i=1}^{n} \log(x_i) - 2n \log(2\lambda) - \sum_{i=1}^{n} \frac{x_i}{\lambda} \left(y_i + \frac{z_i}{2} \right).$$

We don't know the true λ but we do have an estimate and we want to find the expected value so we get

$$\mathbb{E}\left[L(\lambda)|X,Y,Z,\lambda_{k}\right] = 2\sum_{i=1}^{n}\log(x_{i}) - 2n\log(2\lambda) - \mathbb{E}\left[\sum_{i=1}^{n}\frac{x_{i}}{\lambda}\left(y_{i} + \frac{z_{i}}{2}\right)|X,Y,Z,\lambda_{k}\right].$$

The Z data is partially observed so we decompose it into two variables, observed and unobserved, as $Z = \{V, W\}$ where V is the observed part, |V| = r, and W is the unobserved part, |W| = n - r. Then we get

$$\begin{split} \mathbb{E}\left[L(\lambda)|X,Y,V,\lambda_{k}\right] &= 2\sum_{i=1}^{n}\log(x_{i}) - 2n\log(2\lambda) - \sum_{i=1}^{n}\frac{x_{i}}{\lambda}y_{i} - \sum_{i=1}^{r}\frac{x_{i}}{2\lambda}v_{i} - \sum_{i=r+1}^{n}\frac{x_{i}}{2\lambda}\mathbb{E}\left[w_{i}|X,Y,V,\lambda_{k}\right] \\ &= 2\sum_{i=1}^{n}\log(x_{i}) - 2n\log(2\lambda) - \sum_{i=1}^{n}\frac{x_{i}}{\lambda}y_{i} - \sum_{i=1}^{r}\frac{x_{i}}{2\lambda}v_{i} - \sum_{i=r+1}^{n}\frac{x_{i}}{2\lambda}\frac{2\lambda_{k}}{x_{i}} \\ &= 2\sum_{i=1}^{n}\log(x_{i}) - 2n\log(2\lambda) - \sum_{i=1}^{n}\frac{x_{i}}{\lambda}y_{i} - \sum_{i=1}^{r}\frac{x_{i}}{2\lambda}v_{i} - (n-k)\frac{\lambda_{k}}{\lambda} \end{split}$$

To maximize λ we take the derivative and set it to zero and find

$$\lambda = \frac{1}{2n} \left(\sum_{i=1}^{n} x_i y_i + \frac{1}{2} \sum_{i=1}^{r} x_i z_i + (n-r) \lambda_k \right)$$

2.3

```
EM <- function(data, lambdazero, maxiter = 500, eps = 0.001){
    Estep <- function(data,lambda){</pre>
        r <- sum(is.na(data$z))
        n <- nrow(data)</pre>
         (2 * sum(log(data$x)) - 2 * n * log(2 * lambda) -
         sum(data$x * data$y)/lambda -
          sum(data$x * data$z, na.rm = TRUE)/(2*lambda) - (n - r))
    }
    Mstep <- function(data, lambda){</pre>
        n <- nrow(data)</pre>
        r <- sum(is.na(data$z))
        ((sum(data$x * data$y) +
           sum(data$x * data$z, na.rm = TRUE) / 2 +
           (n - r) * lambda) / (2 * n))
    }
    curlambda <- lambdazero
    prevlambda <- lambdazero*5</pre>
    iter <- 1
    while(iter < maxiter && (abs(curlambda - prevlambda) > eps)){
        prevlambda <- curlambda</pre>
        curlambda <- Mstep(data,lambda = curlambda)</pre>
        iter <- iter + 1
    }
    return(list(lambda = curlambda, iter = iter))
}
colnames(physical) <- c("x","y","z")</pre>
res <- EM(data = physical, lambdazero = 100)
print(res)
## $lambda
## [1] 19
##
## $iter
## [1] 16
2.4
lamb <- res$lambda</pre>
physical$EZ <- (2*lamb)/physical$x</pre>
physical$EY <- lamb/physical$x</pre>
```

```
ggplot(physical) +
  geom_point(aes(x=x, y=y), col="blue", alpha = 0.6) +
  geom_point(aes(x=x, y=z), col="red", alpha = 0.6) +
  geom_point(aes(x=x, y=EZ), col="purple", alpha = 0.6) +
  geom_point(aes(x=x, y=EY), col="orange", alpha = 0.6) +
  labs(y = "Y and Z")
```

