Problem Set 4

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Problem 4.2 (g)

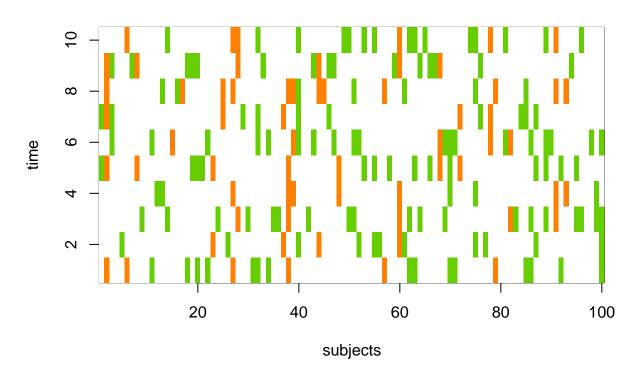
```
data <- read.table("~/Desktop/3_2/6.437/PS/ps4-data/data.txt", quote="\"", comment.char="")
mu = data[, 0:10]
y = data[,11]
mu = as.matrix(mu)
y = as.list(y)</pre>
```

Data Visualization

I color all the exposures of infected subjects(columns) with red. The exposures of uninfected subjects are colored with green.

```
visual = mu
for(i in seq(1, 100, 1)){
   for(j in seq(1, 10, 1)){
      if(y[i]==1 && mu[i, j]==1)
        visual[i, j] = 2
   }
}
cols <- c("#FFFFFF", "chartreuse3", "#FF8000FF")
image(1:nrow(mu), 1:ncol(mu), visual, col=cols, xlab="subjects", ylab="time", main="Disease")</pre>
```

Disease



EM Algorithm

```
T = 5000
z = matrix(0.1, nrow=100, ncol=10)
r = rep(0.5, 10)
for(t in seq(1,T, 1)){
 mur = mu%*%diag(r)
 mur1 = 1 - mur
  prod = matrix(1, 100, 1)
  for(i in seq(1, 10, 1)){
   prod = prod * mur1[ ,i]
  zl = 1 - prod #Denominator of z
  z = diag(as.list(1/zl)) %*% mur
 for(i in seq(1, 100, 1)){
    if(y[i]==0){
     z[i, ] = matrix(0, 1, 10)
    }
 }
 r = colSums(mu * z)/colSums(mu)
print(r)
```

```
## V1 V2 V3 V4 V5
## 4.940656e-324 1.915936e-02 9.881313e-324 2.619160e-01 2.057405e-01
## V6 V7 V8 V9 V10
## 1.208490e-01 1.468435e-01 5.566214e-01 5.622997e-64 1.547256e-01
```