

B565_hw6

2

```
library("quadprog")

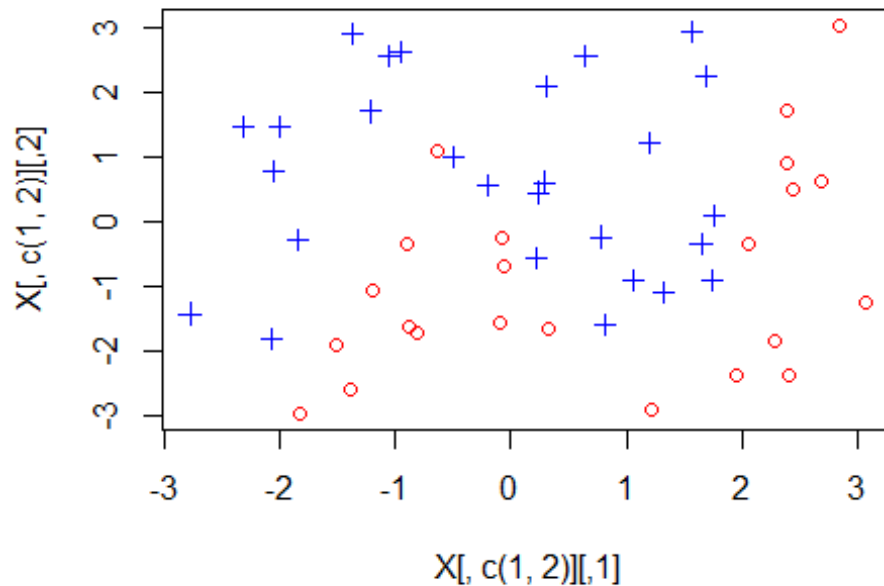
# (a)
# Create w vector
set.seed(100)
w = rnorm(5,0,1)

# Create X of 50X5 matrix, b and Y as class of 50X1 vector
set.seed(100)
X = matrix(0,nrow = 50, ncol = 5)
Y = rep(0,50)

# Update values of X columns
X[,1] = runif(50,min=-3.14,max=3.14)
X[,2] = runif(50,min=-3.14,max=3.14)
X[,3] = cos(X[,1])
X[,4] = sin(X[,1])
X[,5] = rep(1,50)

# Assign sign to Y based on constraints
Y = as.vector(sign(X%%w))

# Plot data X1 with different colours for respective classes
plot(X[,c(1,2)],pch=(Y+2),col=(Y+3),cex=1)
```



```
# (b)
# Solve using quadratic programming

Dmat = diag(5)
Dmat[5,5] = 0.0000001
dvec = rep(0,5)
bvec = rep(1,50)

Amat = as.matrix(X)
A = Amat*Y
result = solve.QP(Dmat,dvec,t(A),bvec)

what = result$solution[1:5]

what

## [1] -37.358591 12.728791 -5.443255 75.626658 10.469340

# Predict for class using result obtained from QP
predict = as.vector(sign(X%%what))
predict

## [1] -1 -1 -1 1 1 -1 -1 -1 1 1 1 -1 1 -1 1 1 1 -1 -1 1 1 1 1
## [24] 1 1 1 1 -1 1 -1 -1 -1 1 -1 -1 -1 1 1 -1 1 1 -1 1 -1 1 -1
## [47] 1 -1 -1 1

Y
```

```
## [1] -1 -1 -1 1 1 -1 -1 -1 1 1 1 -1 1 -1 1 1 1 -1 -1 1 1 1 1
## [24] 1 1 1 1 -1 1 -1 -1 -1 1 -1 -1 -1 1 1 -1 1 1 -1 1 -1 1 -1
## [47] 1 -1 -1 1
```

```
# (c)
```

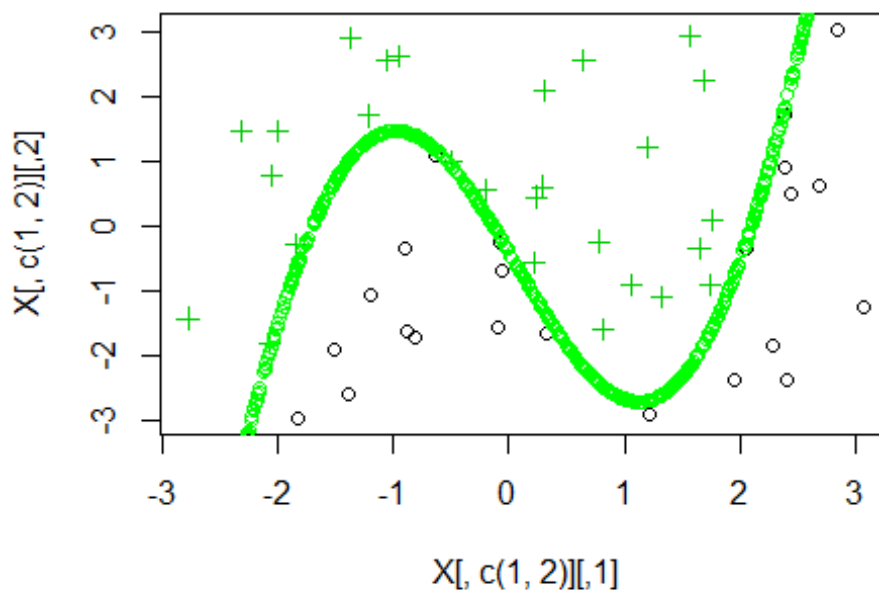
```
# Plot decision boundary which maximize margin
```

```
a = runif(1000,min=-3.14,max=3.14)
```

```
b = (-what[5] - what[1]*a - what[3]*cos(a) - what[4]*sin(a))/what[2]
```

```
plot(X[,c(1,2)],pch=(Y+2),col=(Y+2),cex=1)
```

```
lines(a,b,col="green",type="p")
```



3

```
library("MASS")
```

```
# (a)
```

```
# Create data matrix
```

```
a = scan("D:/Data Mining/hw6/chipotle.dat")
```

```
data = matrix(a,byrow=TRUE,ncol=2)
```

```
data1 = head(data,500)
```

```
# Creating resulting dataframe using cbind
```

```
df = cbind(rep(1,nrow(data1)),data1[,1],data1[,1]^2,data1[,2])
```

```

colnames(df) = c("a", "X1", "X2", "Y")

# X vector
X = df[,c(1,2,3)]

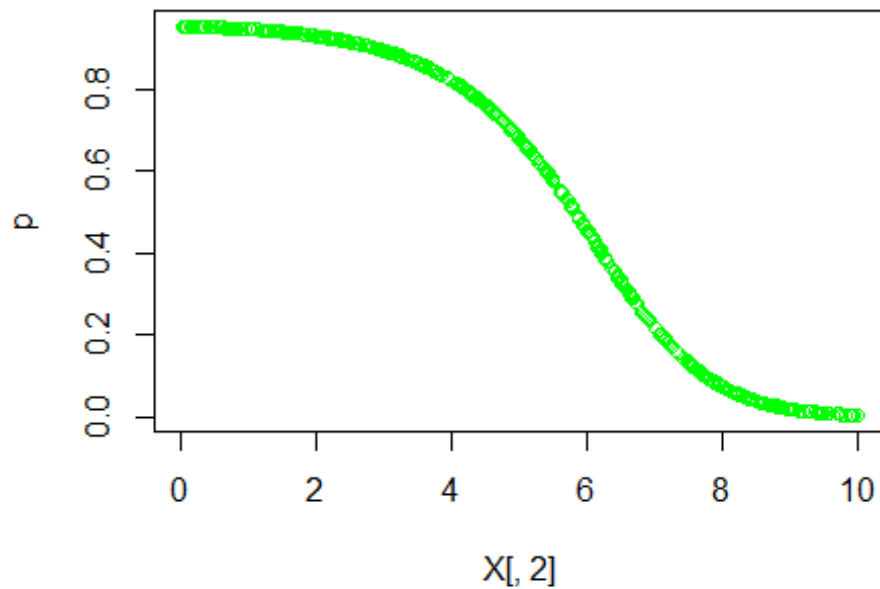
# Initialising w_0

w = matrix(c(3, -.05, -0.08), ncol=1)

# Calculating probability and plotting w.r.t x

p = 1/(1+exp(-t(w) %*% t(X)))
plot(X[,2], p, col="green")

```



```

# (b)

# Gradient descent to optimize w
class = df[,4]
set.seed(100)
w = rep(0,3)
for (i in 1:1000) {
  p = 1/(1+exp(-t(w) %*% t(X)))
  grad = t(X) %*% t(class - p)
  w = w + .01 * grad
  pred = ifelse(p > 0.5, 1, 0)
  accuracy = sum(pred==class)/length(class)
}

```

```

}

w

##           [,1]
## a    37.87352
## X1 229.77711
## X2 -28.88233

# Pred class 1 if p > 0.5 else class 0 and calculate accuracy
pred = ifelse(p > 0.5, 1, 0)
accuracy = sum(pred==class)/length(class)
accuracy

## [1] 0.818

# (c)

# Initialising w
class1 = df[,4]
set.seed(100)
w1 = rep(0,3)

# Newton ralphson method to optimize w
for (i in 1:20) {
  p1 = 1/(1+exp(-t(w1) %*% t(X)))
  p1 = as.vector(p1)
  D = diag(p1*(1-p1))
  H = -t(X) %*% D %*% X
  grad1 = t(X) %*% (class1 - p1)
  w1 = w1 - ginv(H)%*%grad1
}

w1

##           [,1]
## [1,]  2.2067068
## [2,]  0.3801537
## [3,] -0.1259208

# Pred class 1 if p > 0.5 else class 0 and calculate accuracy

pred1 = ifelse(p1 >= 0.5, 1, 0)
accuracy1 = sum(pred1==class1)/length(class1)
accuracy1

## [1] 0.854

```

4

```
# (a)

# Generated 1st cluster points
X1 = matrix(0, nrow = 50, ncol=2)
C1 = rep(1,50)
t1 = matrix(rnorm(2),ncol = 2, nrow = 2)
b1 = matrix(rnorm(2,0,10),ncol = 1, nrow = 2)
for (i in 1:50){
  z1 = matrix(rnorm(2),ncol = 1, nrow = 2)

  X1[i,] = t1%*%z1 + b1
}
X1 = cbind(X1,C1)

# Generated 2nd cluster points
X2 = matrix(0, nrow = 50, ncol=2)
C2 = rep(2,50)
t2 = matrix(rnorm(2),ncol = 2, nrow = 2)
b2 = matrix(rnorm(2,0,10),ncol = 1, nrow = 2)
for (i in 1:50){
  z2 = matrix(rnorm(2),ncol = 1, nrow = 2)

  X2[i,] = t2%*%z2 + b2
}

X2 = cbind(X2,C2)

# Generated 3rd cluster points
X3 = matrix(0, nrow = 50, ncol=2)
C3 = rep(3,50)
t3 = matrix(rnorm(2),ncol = 2, nrow = 2)
b3 = matrix(rnorm(2,0,10),ncol = 1, nrow = 2)
for (i in 1:50){
  z3 = matrix(rnorm(2),ncol = 1, nrow = 2)

  X3[i,] = t3%*%z3 + b3
}
X3 = cbind(X3,C3)

# Generated 4th cluster points
X4 = matrix(0, nrow = 50, ncol=2)
C4 = rep(4,50)
t4 = matrix(rnorm(2),ncol = 2, nrow = 2)
b4 = matrix(rnorm(2,0,10),ncol = 1, nrow = 2)
for (i in 1:50){
```

```

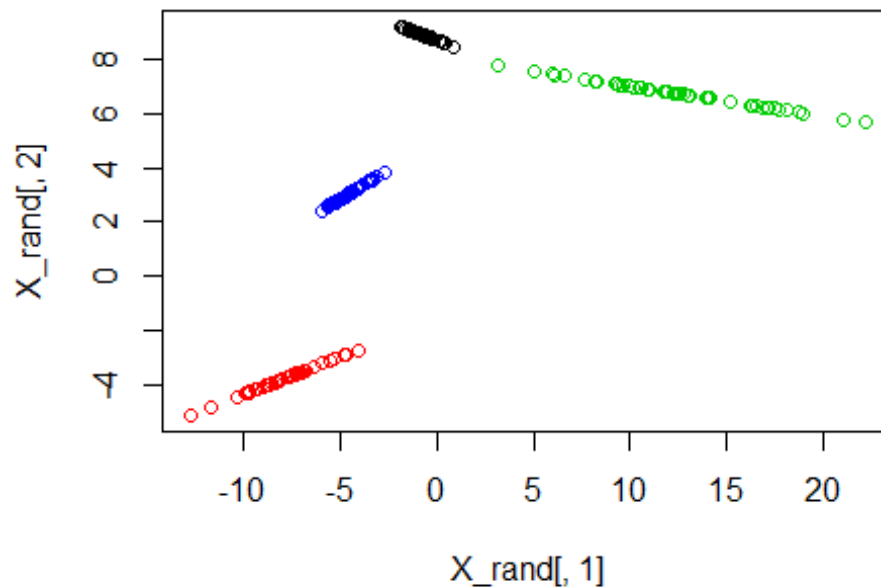
z4 = matrix(rnorm(2),ncol = 1, nrow = 2)

X4[i,] = t4**z4 + b4
}
X4 = cbind(X4,C4)

# Shuffling data for non-biased initialisation of prototype points
X = rbind(X1,X2,X3,X4)
X_rand = X[sample(nrow(X)),]

# Plotting data points of different cluster with different colour
plot(X_rand[,1],X_rand[,2],col=X_rand[,3])

```



```

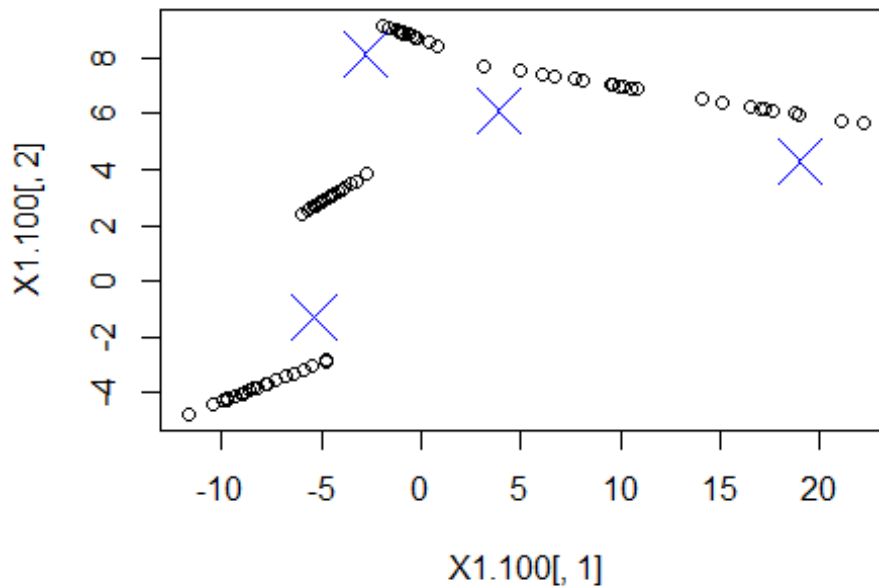
# (c)

X.100 = X[sample(nrow(X),100),]

K = 4
X1.100 = cbind(X.100[,c(1,2)],rep(1,100))
#m1 = X1.100[sample(nrow(X1.100),4),c(1,2)]
m1 = matrix(0,nrow=4,ncol=2)
m1[,1] = runif(4,min=min(X1.100[,1]),max=max(X1.100[,1]))
m1[,2] = runif(4,min=min(X1.100[,2]),max=max(X1.100[,2]))

```

```
# Plotting the scatter plot with initial cluster same for all points
plot(X1.100[,1], X1.100[,2],col=X1.100[,3])
lines(m1[,1],m1[,2],col="blue",pch=4,cex=3,type="p")
```

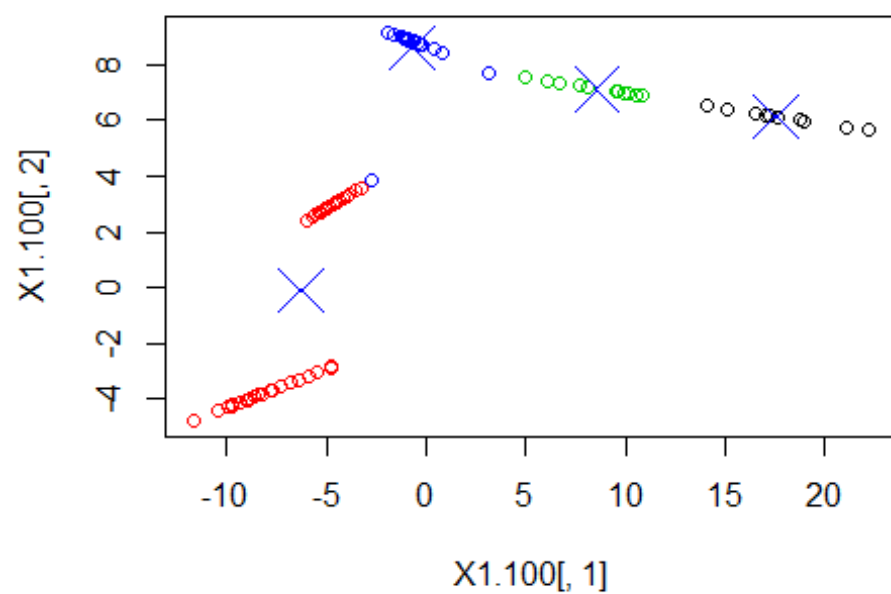
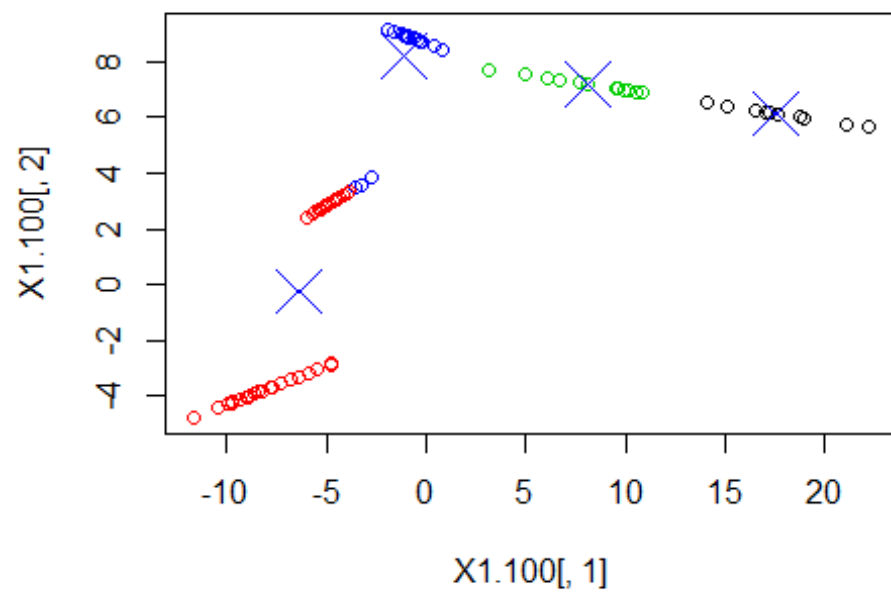


```
# Looping through 50 iteration with user input of cluster formation
dist_matrix1 = matrix(0,nrow=100,ncol= 4)

H = c()
steps = c()
mnew = matrix(0,ncol=2,nrow=4)
for (i in 1:100){
  for (j in 1:nrow(m1)) {
    dist_matrix1[,j] = sqrt((X1.100[,1]-m1[j,1])^2+(X1.100[,2]-m1[j,2])^2)
  }
  cluster = apply(dist_matrix1, 1, which.min)
  X1.100[,3] = cluster
  for (k in 1:nrow(m1)){
    subset = X1.100[X1.100[,3] == k,]
    m1[k,1] = mean(subset[,1])
    m1[k,2] = mean(subset[,2])
  }
}
H1 = sum(apply(dist_matrix1, 1, min))
plot(X1.100[,1], X1.100[,2],col=X1.100[,3])
lines(m1[,1],m1[,2],col="blue",pch=4,cex=3,type="p")
```

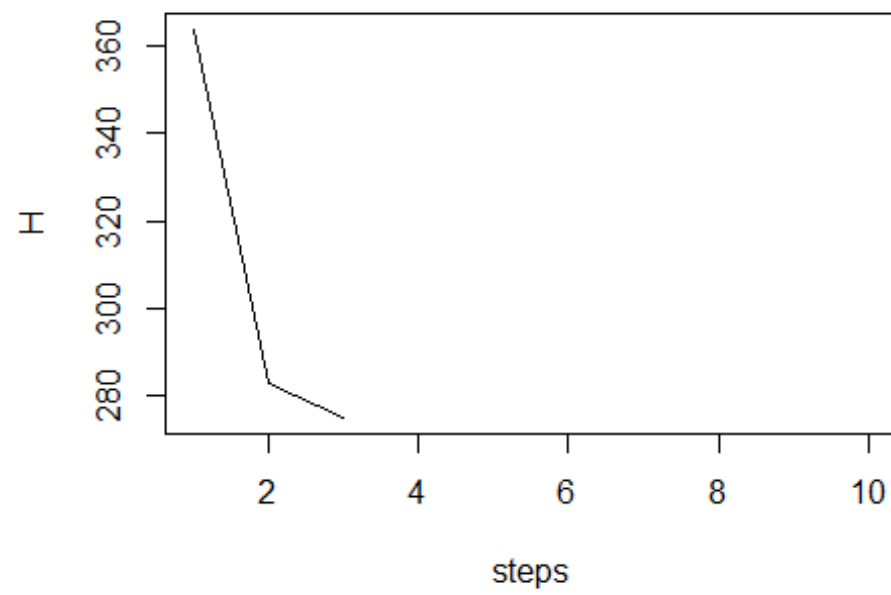


```
if (all(m1 == mnew) == TRUE){  
  H = c(H,H1)  
  steps = c(steps,i)  
  K = i  
  cat("Number of steps to converge: ", K)  
  break  
} else {  
  H = c(H,H1)  
  steps = c(steps,i)  
  mnew = m1  
  
}  
}
```



```
## Number of steps to converge: 3
```

```
plot(steps,H,type="l",xlim = c(1,10))
```



K

[1] 3