

Homework 1

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January 22, 2020

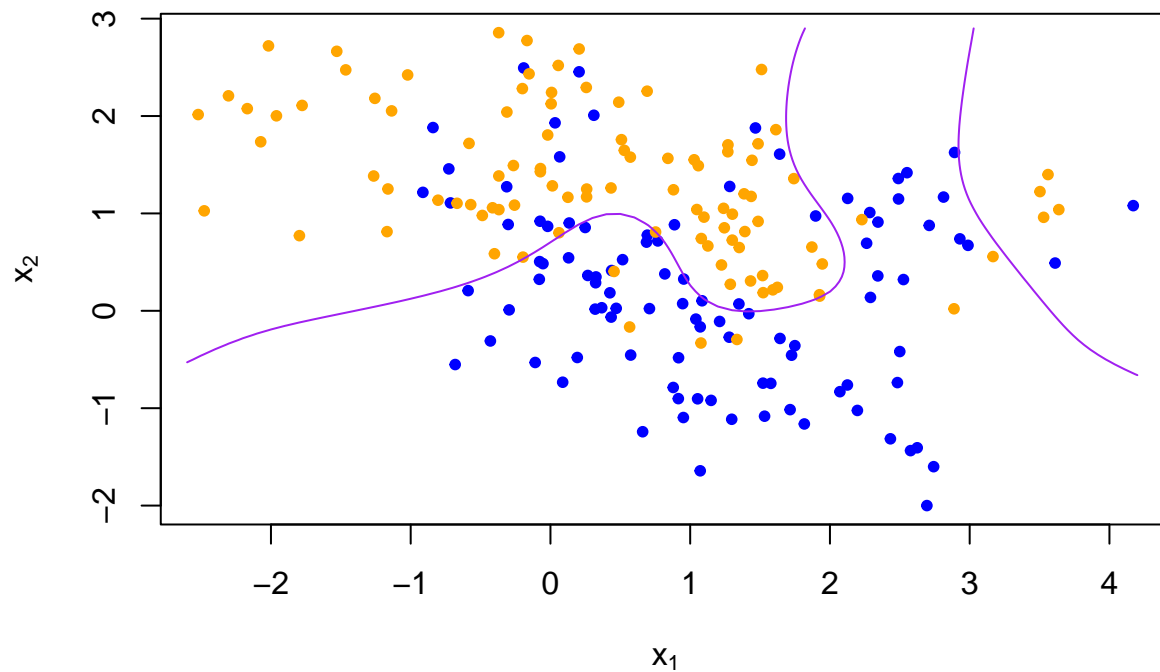
Rewrite code with lm()

```
library('class')
library('dplyr')

##
## Attaching package: 'dplyr'
## The following objects are masked from 'package:stats':
##
##   filter, lag
## The following objects are masked from 'package:base':
##
##   intersect, setdiff, setequal, union
## load binary classification example data from author website
## 'ElemStatLearn' package no longer available
load(url('https://web.stanford.edu/~hastie/ElemStatLearn/datasets/ESL.mixture.rda'))
dat <- ESL.mixture

plot_mix_data <- expression({
  plot(dat$x[,1], dat$x[,2],
       col=ifelse(dat$y==0, 'blue', 'orange'),
       pch=20,
       xlab=expression(x[1]),
       ylab=expression(x[2]))
  ## draw Bayes (True) classification boundary
  prob <- matrix(dat$prob, length(dat$px1), length(dat$px2))
  cont <- contourLines(dat$px1, dat$px2, prob, levels=0.5)
  rslt <- sapply(cont, lines, col='purple')
})

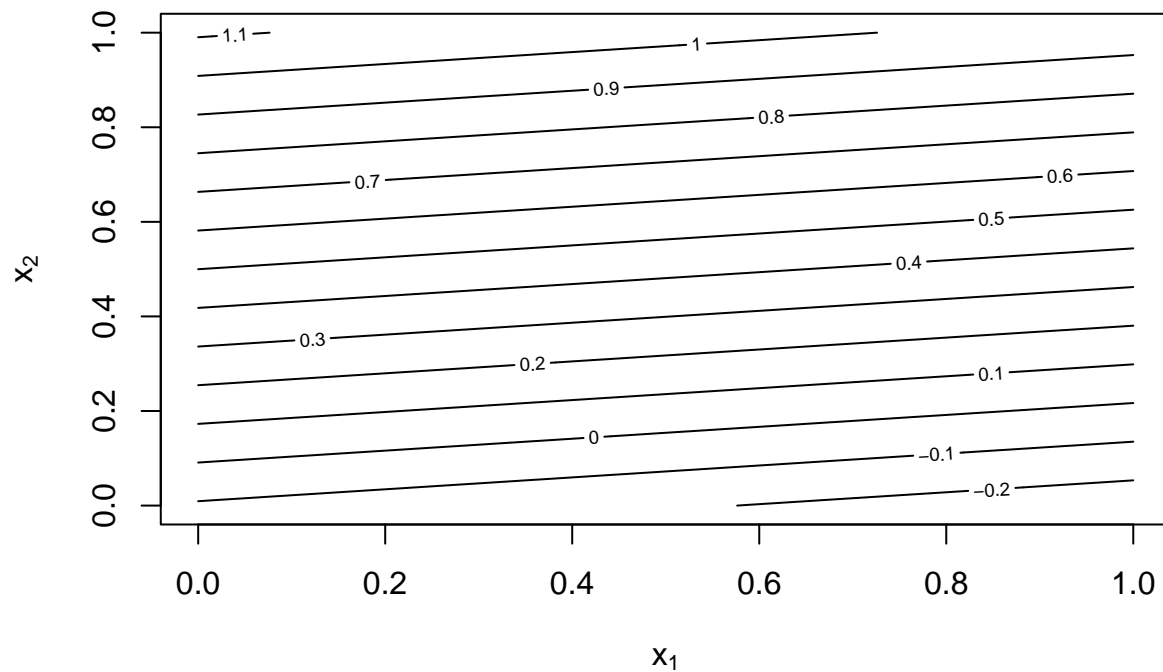
eval(plot_mix_data)
```



```
## fit linear classifier
fit_lc <- function(y,x){
  return(lm(y~x+1))
}

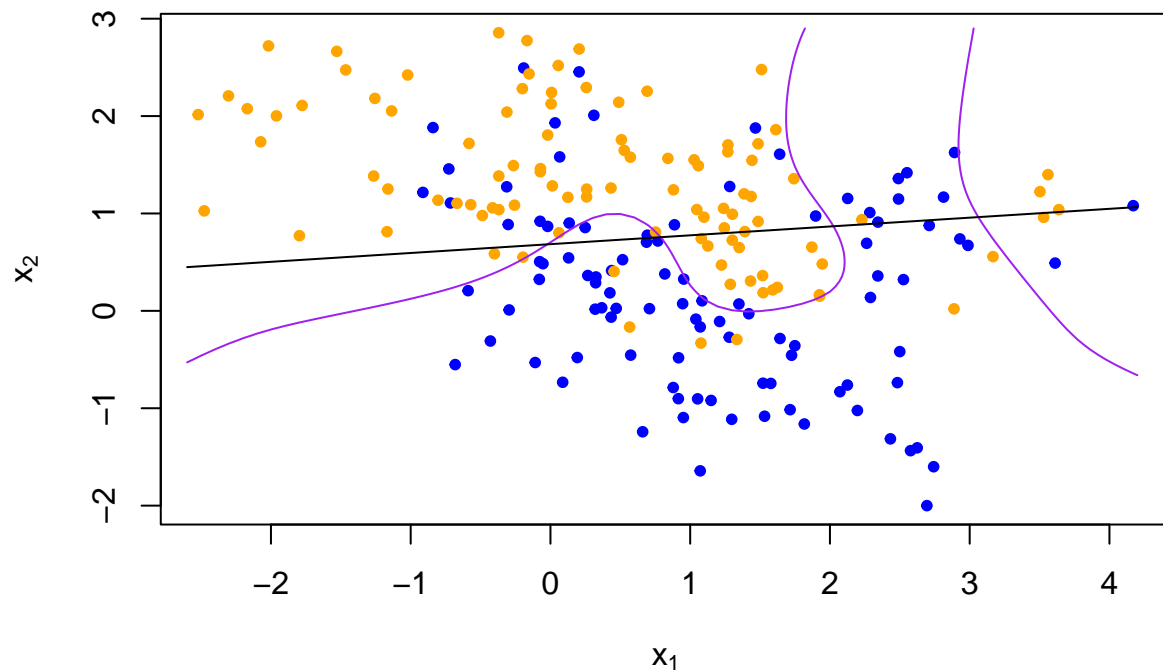
## make predictions from linear classifier
predict_lc <- function(x, fun) {
  cbind(1, x) %*% fun$coefficients
}

## fit model to mixture data and make predictions
lc_beta <- fit_lc(dat$y, dat$x)
lc_pred <- predict_lc(dat$xnew, lc_beta)
## reshape predictions as a matrix
lc_pred <- matrix(lc_pred, length(dat$px1), length(dat$px2))
contour(lc_pred,
  xlab=expression(x[1]),
  ylab=expression(x[2]))
```



```
## find the contours in 2D space such that lc_pred == 0.5
lc_cont <- contourLines(dat$px1, dat$px2, lc_pred, levels=0.5)

## plot data and decision surface
eval(plot_mix_data)
sapply(lc_cont, lines)
```



```
## [[1]]
## NULL

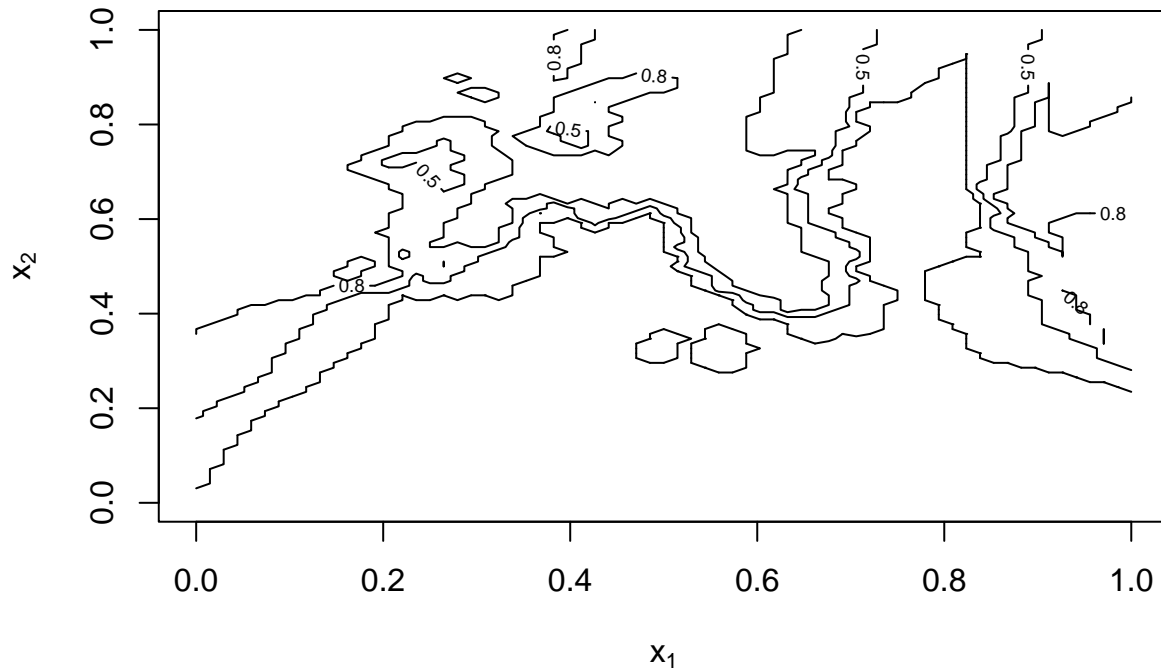
## fit knn classifier
## use 5-NN to estimate probability of class assignment
```

```

knn_fit <- knn(train=dat$x, test=dat$xnew, cl=dat$y, k=5, prob=TRUE)
knn_pred <- attr(knn_fit, 'prob')
knn_pred <- ifelse(knn_fit == 1, knn_pred, 1-knn_pred)

## reshape predictions as a matrix
knn_pred <- matrix(knn_pred, length(dat$px1), length(dat$px2))
contour(knn_pred,
        xlab=expression(x[1]),
        ylab=expression(x[2]),
        levels=c(0.2, 0.5, 0.8))

```

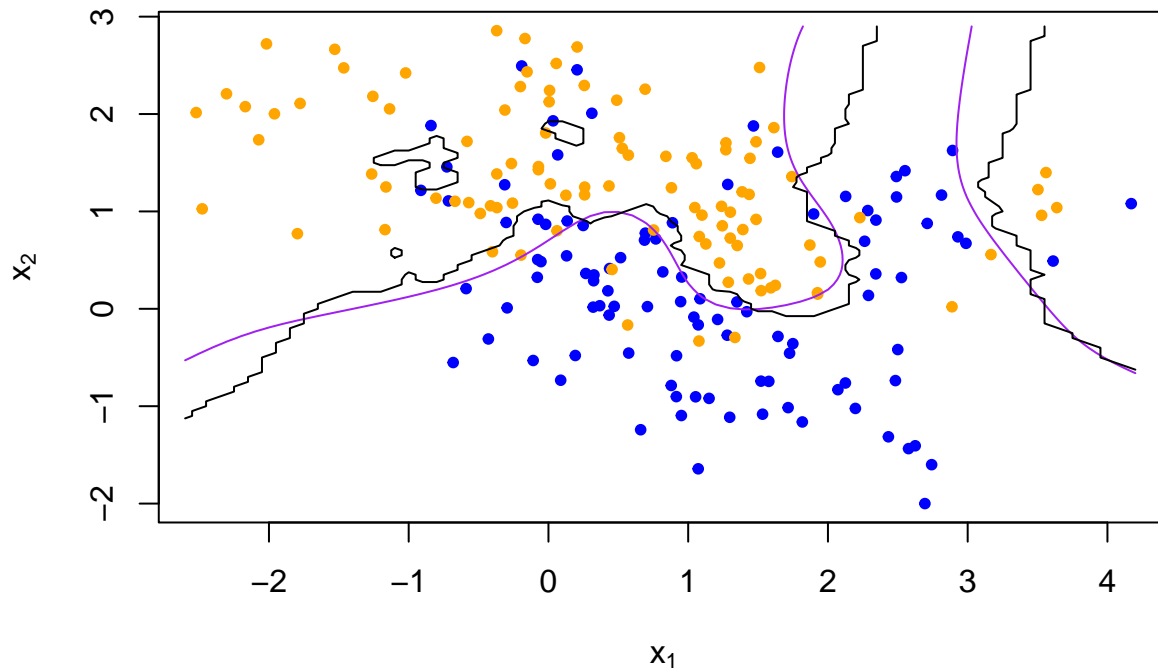


```

## find the contours in 2D space such that knn_pred == 0.5
knn_cont <- contourLines(dat$px1, dat$px2, knn_pred, levels=0.5)

## plot data and decision surface
eval(plot_mix_data)
sapply(knn_cont, lines)

```



```
## [[1]]
## NULL
##
## [[2]]
## NULL
##
## [[3]]
## NULL
##
## [[4]]
## NULL
##
## [[5]]
## NULL

## do bootstrap to get a sense of variance in decision surface
resample <- function(dat) {
  idx <- sample(1:length(dat$y), replace = T)
  dat$y <- dat$y[idx]
  dat$x <- dat$x[idx,]
  return(dat)
}

## plot linear classifier for three bootstraps
par(mfrow=c(1,3))
for(b in 1:3) {
  datb <- resample(dat)
  ## fit model to mixture data and make predictions
  lc_beta <- fit_lc(datb$y, datb$x)
  lc_pred <- predict_lc(datb$xnew, lc_beta)

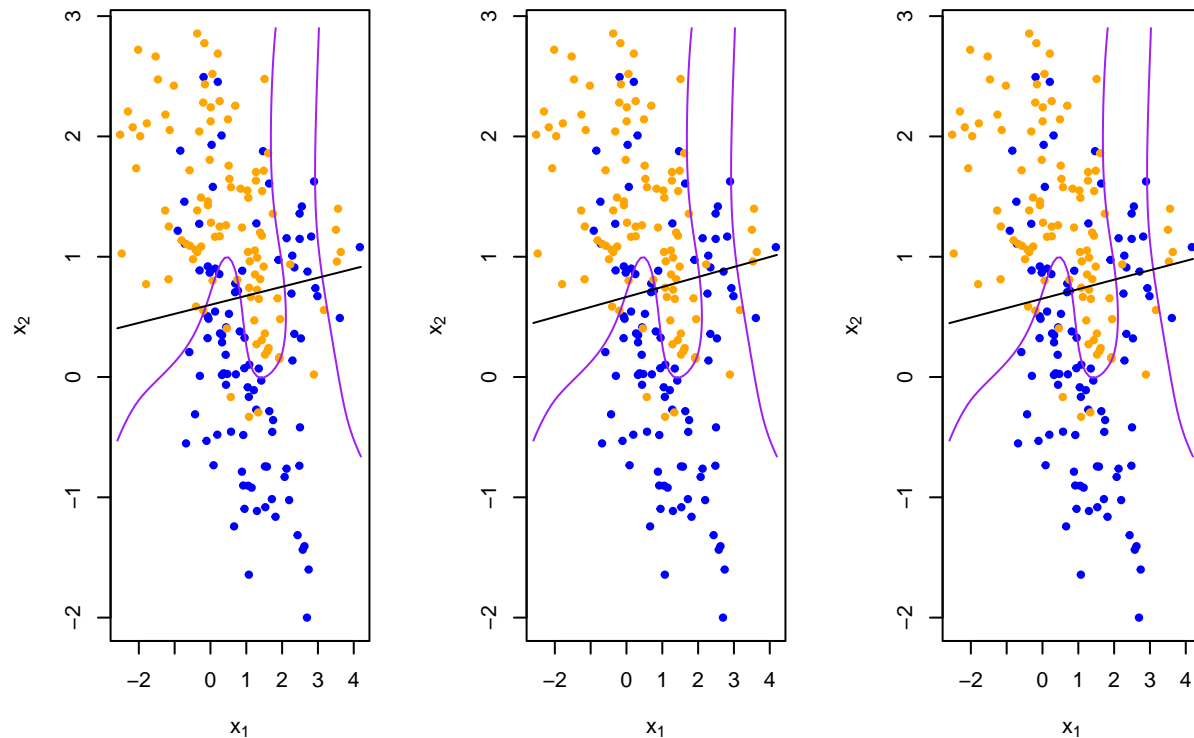
  ## reshape predictions as a matrix
  lc_pred <- matrix(lc_pred, length(datb$px1), length(datb$px2))
}
```

```

## find the contours in 2D space such that lc_pred == 0.5
lc_cont <- contourLines(datb$px1, datb$px2, lc_pred, levels=0.5)

## plot data and decision surface
eval(plot_mix_data)
sapply(lc_cont, lines)
}

```



```

## plot 5-NN classifier for three bootstraps
par(mfrow=c(1,3))
for(b in 1:3) {
  datb <- resample(dat)

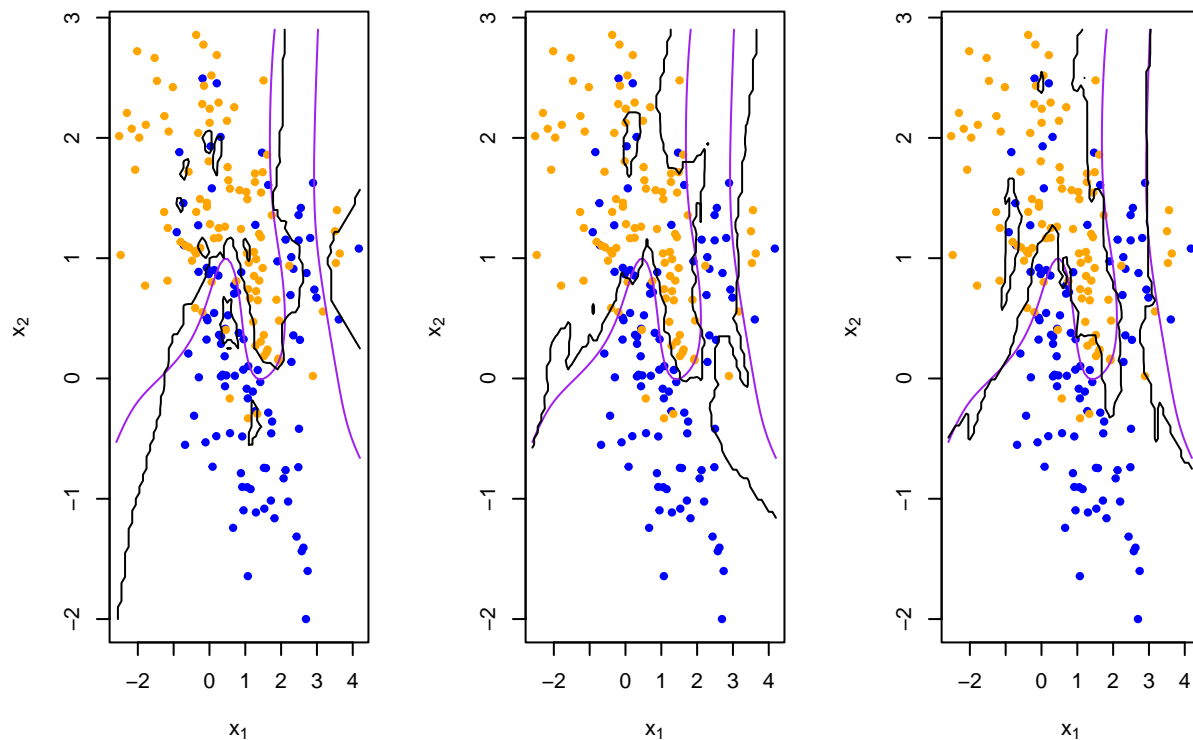
  knn_fit <- knn(train=datb$x, test=datb$xnew, cl=datb$y, k=5, prob=TRUE)
  knn_pred <- attr(knn_fit, 'prob')
  knn_pred <- ifelse(knn_fit == 1, knn_pred, 1-knn_pred)

  ## reshape predictions as a matrix
  knn_pred <- matrix(knn_pred, length(datb$px1), length(datb$px2))

  ## find the contours in 2D space such that knn_pred == 0.5
  knn_cont <- contourLines(datb$px1, datb$px2, knn_pred, levels=0.5)

  ## plot data and decision surface
  eval(plot_mix_data)
  sapply(knn_cont, lines)
}

```



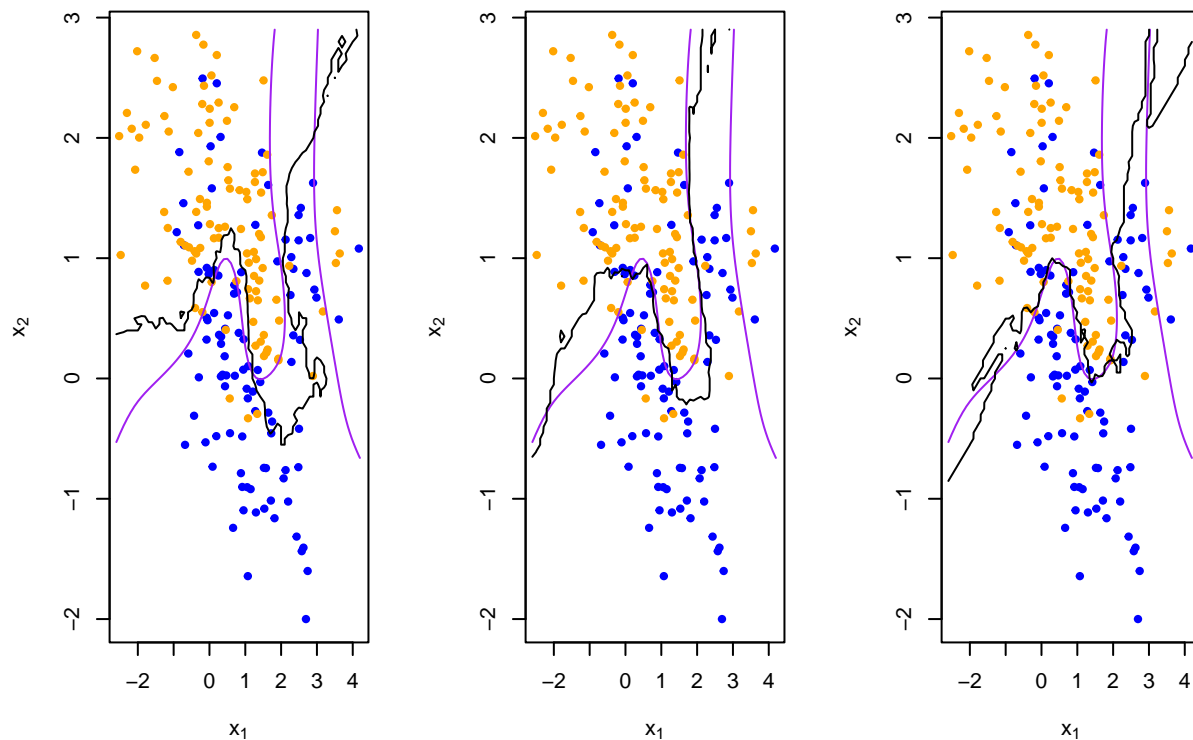
```
## plot 20-NN classifier for three bootstraps
par(mfrow=c(1,3))
for(b in 1:3) {
  datb <- resample(dat)

  knn_fit <- knn(train=datb$x, test=datb$xnew, cl=datb$y, k=20, prob=TRUE)
  knn_pred <- attr(knn_fit, 'prob')
  knn_pred <- ifelse(knn_fit == 1, knn_pred, 1-knn_pred)

  ## reshape predictions as a matrix
  knn_pred <- matrix(knn_pred, length(datb$px1), length(datb$px2))

  ## find the contours in 2D space such that knn_pred == 0.5
  knn_cont <- contourLines(datb$px1, datb$px2, knn_pred, levels=0.5)

  ## plot data and decision surface
  eval(plot_mix_data)
  sapply(knn_cont, lines)
}
```



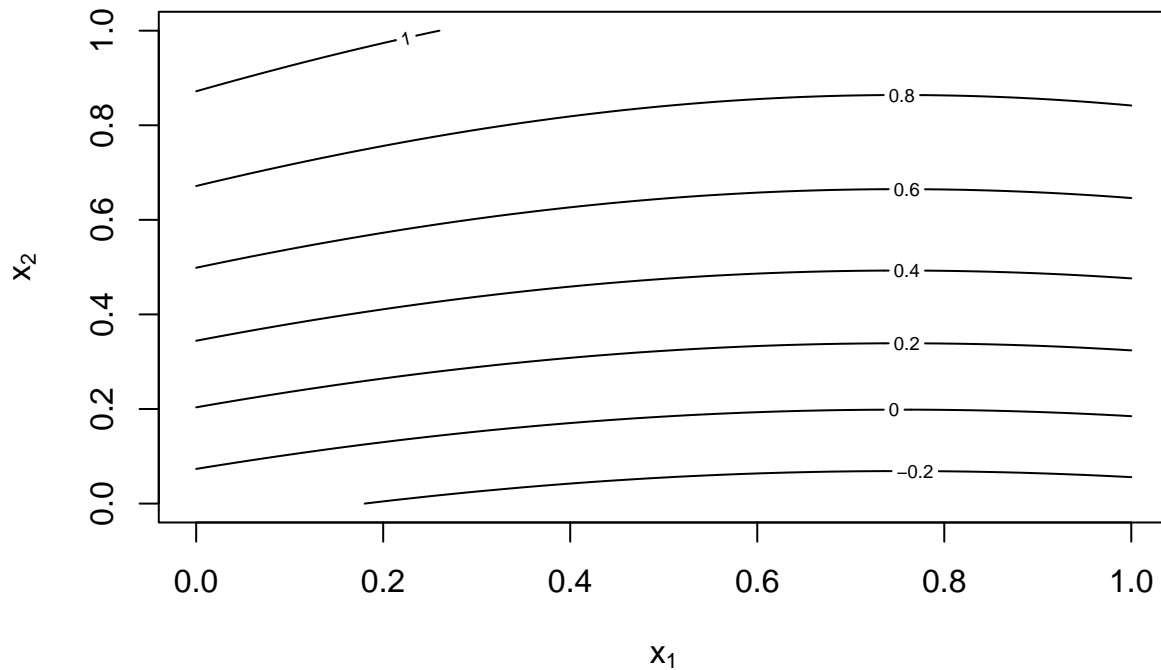
Use squared x to fit the data

```
## fit linear classifier
fit_lc <- function(y,x){

  return(lm(y~I(x^2) + I(x)+1))
}

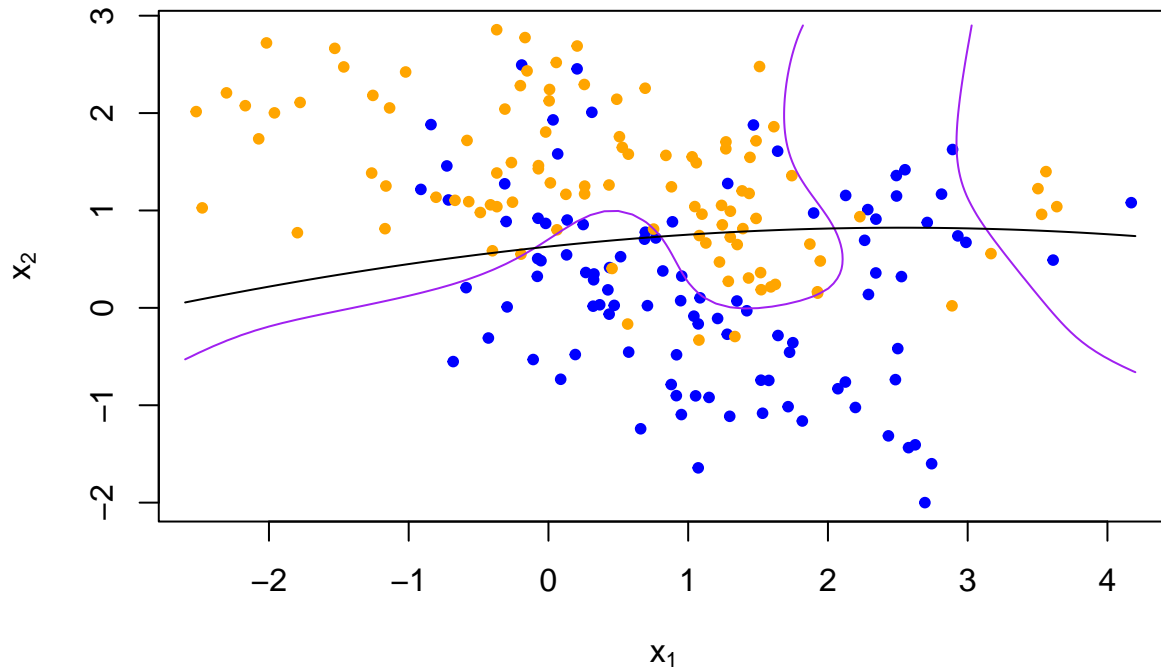
## make predictions from linear classifier
predict_lc <- function(x, fun) {
  cbind(cbind(1, x*x),x) %*% fun$coefficients
}

## fit model to mixture data and make predictions
lc_beta_new <- fit_lc(dat$y, dat$x)
lc_pred_new <- predict_lc(dat$xnew, lc_beta_new)
## reshape predictions as a matrix
lc_pred_new <- matrix(lc_pred_new, length(dat$px1), length(dat$px2))
contour(lc_pred_new,
        xlab=expression(x[1]),
        ylab=expression(x[2]))
```

```
## find the contours in 2D space such that lc_pred == 0.5
lc_cont_new <- contourLines(dat$px1, dat$px2, lc_pred_new, levels=0.5)

## plot data and decision surface
eval(plot_mix_data)
sapply(lc_cont_new, lines)
```



```
## [[1]]
## NULL

## do bootstrap to get a sense of variance in decision surface
resample <- function(dat) {
```

```

idx <- sample(1:length(dat$y), replace = T)
dat$y <- dat$y[idx]
dat$x <- dat$x[idx,]
return(dat)
}

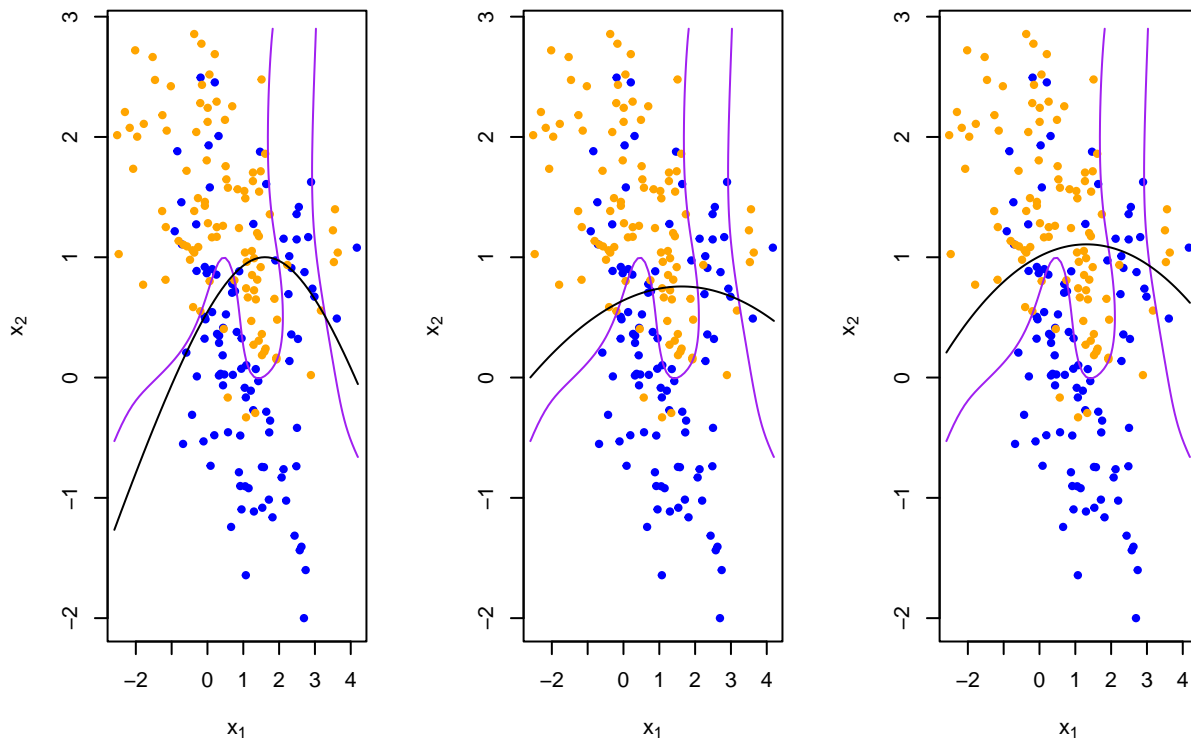
## plot linear classifier for three bootstraps
par(mfrow=c(1,3))
for(b in 1:3) {
  datb <- resample(dat)
  ## fit model to mixture data and make predictions
  lc_beta_new <- fit_lc(datb$y, datb$x)
  lc_pred_new <- predict_lc(datb$xnew, lc_beta_new)

  ## reshape predictions as a matrix
  lc_pred_new <- matrix(lc_pred_new, length(datb$px1), length(datb$px2))

  ## find the contours in 2D space such that lc_pred == 0.5
  lc_cont_new <- contourLines(datb$px1, datb$px2, lc_pred_new, levels=0.5)

  ## plot data and decision surface
  eval(plot_mix_data)
  sapply(lc_cont_new, lines)
}

```



```
var(as.vector(lc_pred))
```

```
## [1] 0.1250923
```

```
var(as.vector(lc_pred_new))
```

```
## [1] 0.1551389
result1 <- as.vector(lc_pred)
result2 <- as.vector(lc_pred_new)
p1<-rep(0,1)
p2<-rep(0,1)
for (i in 1:length(result1)){
  if ((result1[i] >= 0.5 & dat$marginal[i]>= 0.5)|(result1[i] <= 0.5 & dat$marginal[i]<= 0.5)){
    p1[i] <- 0
  }
  else{
    p1[i] <- 1
  }

  if ((result2[i] >= 0.5 & dat$marginal[i]>= 0.5)|(result2[i] <= 0.5 & dat$marginal[i]<= 0.5)){
    p2[i] <- 0
  }
  else{
    p2[i] <- 1
  }
}
mean(p1)

## [1] 0.4464939
mean(p2)
```

```
## [1] 0.4169229
```

From the result we can see that the variance of the fit model which has squared x has smaller bias and variance