

Simulated ROC

jdt

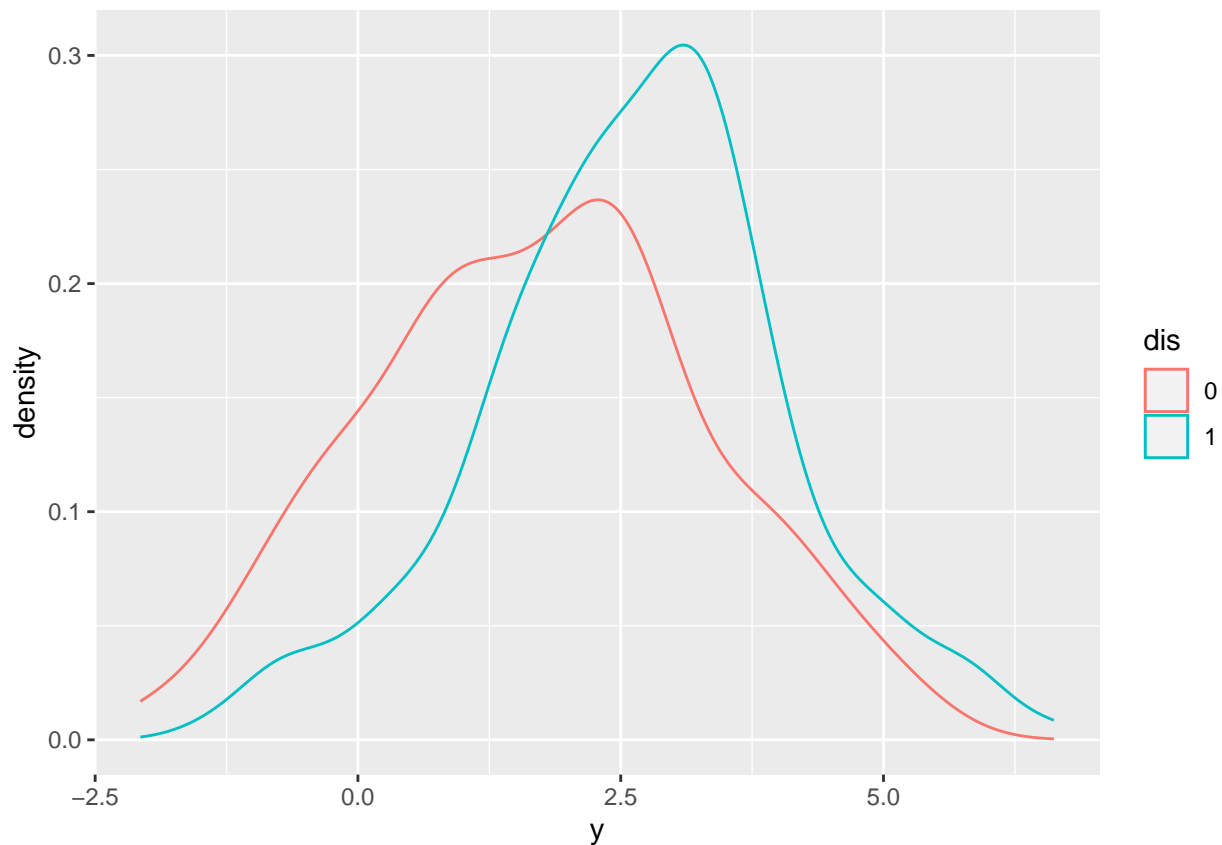
12/13/2020

Set seed for the simulation

```
# clear the environment and set seed  
rm(list = ls())  
set.seed(12345)
```

Function to generate normal data. For dis=0 (“Control”) and dis=1 (“Disease”). The separation between the two groups is controlled by one’s choice of c0, c1, sd_e0, and sd_e1. n0 and n1 are the sample sizes for the groups.

```
library(ggplot2)  
  
gen_Norm_data = function(c0,sd_e0,n0,c1,sd_e1,n1){  
  eps0 = rnorm(n0,0,sd_e0)  
  eps1 = rnorm(n1,0,sd_e1)  
  y0 = c0 + eps0  
  y1 = c1 + eps1  
  
  data.frame(y = c(y0, y1),  
             dis = as.factor(c(rep(0,n0), rep(1,n1))))  
}  
dat1 = gen_Norm_data(1.5,1.5,200,2.5,1.5,200)  
  
#plot density functions  
ggplot(dat1, aes(color = dis, y)) + geom_density()
```



Create discrete variables for Y

```
y=dat1[,1]
summary(y)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
## -2.071  1.084   2.296   2.172  3.225   6.621
```

```
case=dat1[,2]
```

Create binary table with cutoff $y > 2.5$

```
high_y = y > 2.5
tex1 = table(case,high_y)
addmargins(tex1)
```

```
##      high_y
## case FALSE TRUE Sum
##  0     137   63 200
##  1      87  113 200
##  Sum    224  176 400
```

```
prop.out = prop.table(tex1,1)
specificity = prop.out[1,1]
sensitivity = prop.out[2,2]
prop.out
```

```
##      high_y
## case FALSE TRUE
##  0 0.685 0.315
##  1 0.435 0.565
```

```
sensitivity
```

```
## [1] 0.565
```

```
specificity
```

```
## [1] 0.685
```

```
TPR = sensitivity
```

```
FPR = 1 - specificity
```

```
TPR
```

```
## [1] 0.565
```

```
FPR
```

```
## [1] 0.315
```

Create binary table with cutoff $y > 3.0$

```
high_y = y > 3.0
```

```
tex1 = table(case,high_y)
```

```
addmargins(tex1)
```

```
##      high_y
```

```
## case FALSE TRUE Sum
```

```
##  0      161   39 200
```

```
##  1      115   85 200
```

```
## Sum      276  124 400
```

```
prop.out = prop.table(tex1,1)
```

```
specificity = prop.out[1,1]
```

```
sensitivity = prop.out[2,2]
```

```
prop.out
```

```
##      high_y
```

```
## case FALSE  TRUE
```

```
##  0 0.805 0.195
```

```
##  1 0.575 0.425
```

```
sensitivity
```

```
## [1] 0.425
```

```
specificity
```

```
## [1] 0.805
```

```
TPR = sensitivity
```

```
FPR = 1 - specificity
```

```
TPR
```

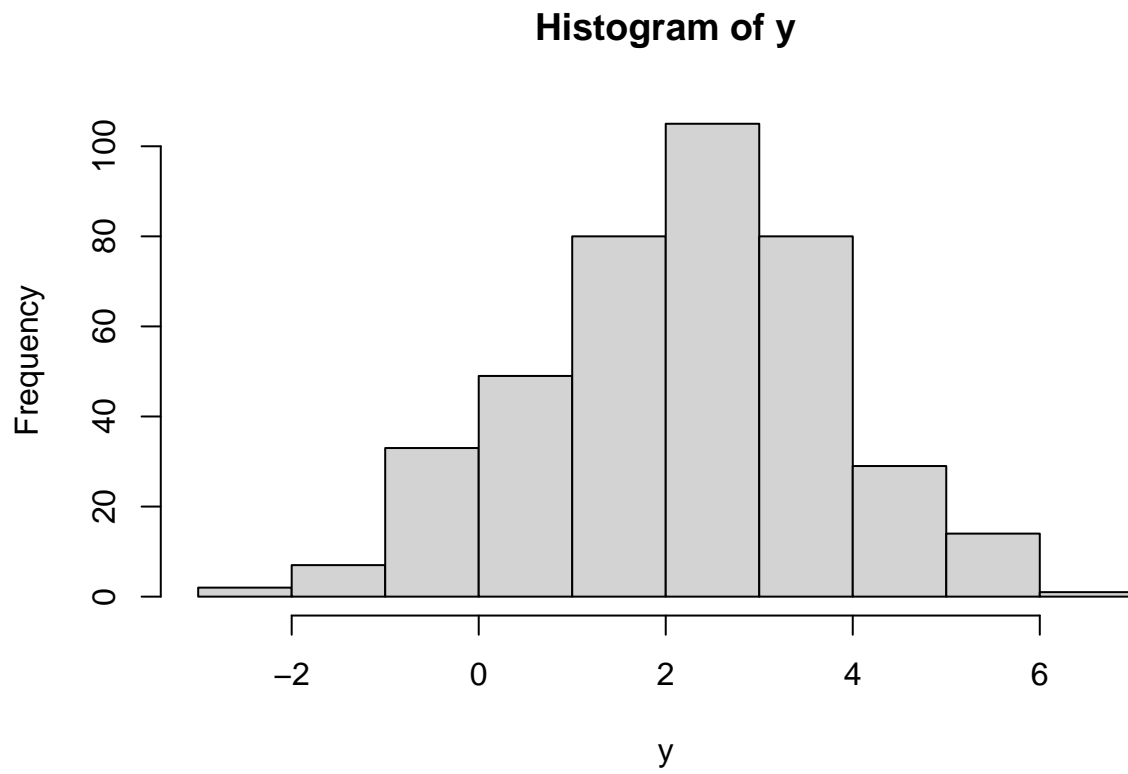
```
## [1] 0.425
```

```
FPR
```

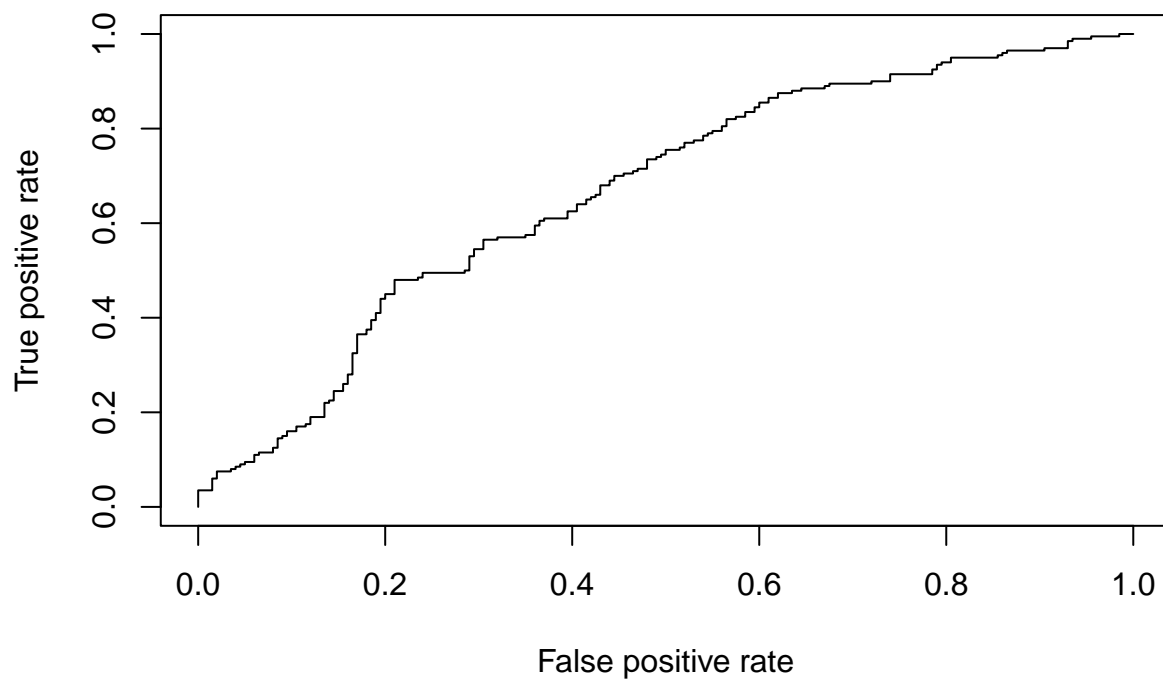
```
## [1] 0.195
```

Instead of creating a table at each cutoff point one can construct a ROC plot for a continuous variable Y. Which is constructing using the pairs (TPR, FPR) at each cutoff point (in R). Sometimes (as in SAS) this curve is smoothed.

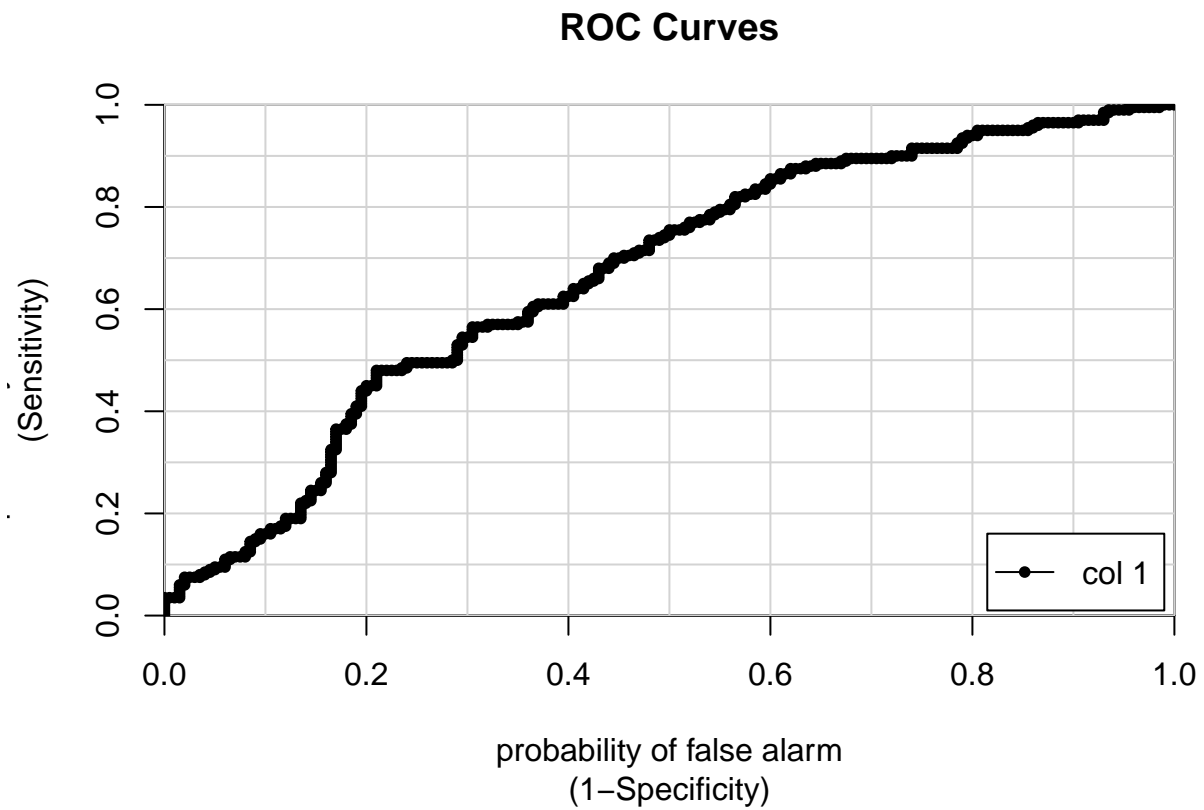
```
library(ROCR)
hist(y)
```



```
pred = prediction(y, case)
perf=performance(pred, "tpr", "fpr")
plot(perf)
```



```
library(caTools)
colAUC(y, case, plotROC=TRUE)
```



```
##           [,1]
## 0 vs. 1 0.66575
```