# LDA: Introduction

#### Introduction

Linear Disriminant Analysis (LDA) is method of making classifications based on an underlying assumption of normality as a means of approximating the Bayes Classifier.

The idea is based on Bayes Rule.

$$P(Y = k|X = x) = \frac{P(X = x|Y = k)P(Y = k)}{P(X = x)}$$

The denominator isn't actually relevant since it is indepedent of the class k. In any case, it's given by the "law of total probability.," namely

$$P(X = x) = \sum_{k=1}^{C} P(X = x | Y = k) P(Y = k)$$

For any input value X = x, the Bayes Classifier simply picks the class with the highest probability. Formally,

$$f(x) = \operatorname{argmax}_k P(Y = k | X = x)$$

To use this approach in practice, we need to estimate the various quantites that go into Bayes Rule. To start, we assume that, given the class k, that the values x are normally distributed with a mean  $\mu_k$  and a comman variance  $\sigma^2$ . In other words,

$$P(X = x|Y = k) = \frac{1}{\sqrt{2\pi}\sigma}e^{-(x-\mu_k)^2/(2\sigma^2)}$$

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Use  $\pi_k = P(Y = k)$ , the probability that an observation. is in class k.

That being said, we get:

$$P(Y = k|X = x) \propto \frac{1}{\sqrt{2\pi}\sigma} e^{-(x-\mu_k)^2/(2\sigma^2)} \times \pi_k$$

It's easier to work with the log of this expression.

$$\log(P(Y=k|X=x)) = CONSTANT - \sqrt{2\pi} - \sigma - (x - \mu_k)^2/(2\sigma^2) + \log(\pi_k)$$

The CONSTANT comes from the denominator (independent of k).

This is the discriminant function for the class k, write it as

$$f_k(x) = CONSTANT - \sqrt{2\pi} - \sigma - (x - \mu_k)^2 / (2\sigma^2) + \log(\pi_k)$$

As before, we classify x into the class k for which this function is largest. This means we can ignore any additive terms which don't involve k. After multiplying out the square and eliminating unneeded terms, we get a simpler expression.

$$f_k(x) = x\frac{\mu_k}{\sigma^2} - \frac{\mu_k}{2\sigma^2} + \log(\pi_k)$$

## Parameter Estimates

Now we need good estimates of the values  $\pi_k$ ,  $\mu_k$  and  $\sigma^2$  for k = 1, 2, ..., C where C = the number of classes.

The first easy,  $\pi_k$  is just observed the class proportion. That is, if  $N_k$  is the number of observations in class k and N, then

$$\pi_k \approx \frac{N_k}{N}.$$

Next,  $\mu_k$  is just the mean of x in class k.

An estimate of  $\sigma^2$  is a bit deeper (but not a lot deeper).

data.df <- read\_csv(file.path(dataDir,file))</pre>

$$\sigma^{2} = \frac{1}{N - C} \sum_{k=1}^{C} (N_{k} - 1) \sigma_{k}^{2}$$

where  $\sigma_k^2$  is the variance of x in class k.

Given a data frame, all of these quantities are easily computed in R.

#### **Example Data**

Load the libraries.

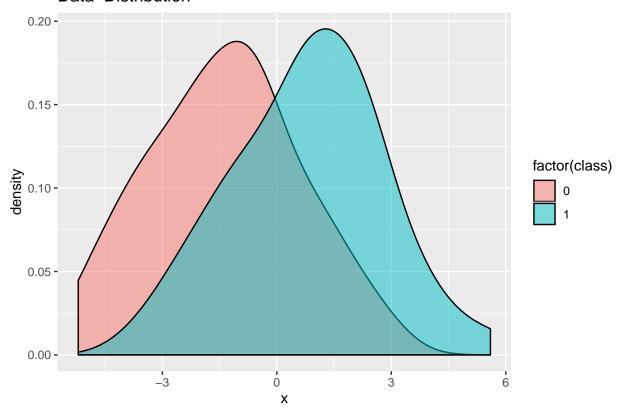
```
library(tidyverse)
## -- Attaching packages ----- tidyverse 1.2.1 --
                                      0.3.3
## v ggplot2 3.2.1
                       v purrr
## v tibble 2.1.3 v dplyr 0.8.4
## v tidyr 1.0.2 v stringr 1.4.0
## v readr 1.3.1 v forcats 0.3.0
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                        masks stats::lag()
library(MASS) ##For lda
## Attaching package: 'MASS'
## The following object is masked from 'package:dplyr':
##
##
        select
Here is a simple data set with p = 1 and C = 2.
dataDir <- "DATA"</pre>
file <- "LDA_ClassData1.csv"</pre>
```

```
## Parsed with column specification:
## cols(
##    x = col_double(),
##    class = col_double()
## )
```

Note: The variable class is categorical. You might want to mutate it to a factor.

```
data.df %>%
   ggplot()+
   geom_density(aes(x,fill=factor(class)),alpha=0.5)+
   labs(title="Data Distribution")
```

## Data Distribution



Peek at the data...how many classes? What are the counts, means and variances of each class?

```
with(data.df,table(class))
## class
## 0 1
```

### Build your own Linear Discriminant Analysis classifier

## 56 44

Here you build your own discriminant functions directly from the data. Everything follows the theory above.

Compute the mean and variances of each class. Take a look at this data to get an idea.

Plan: Separate out two data frames: data0.df (only class == ) and data1.df (class==1)

```
data0 <- data.df %>%
  filter(class==0)
data1 <- data.df %>%
  filter(class==1)
```

Extract all the important parameters.  $n_0$ ,  $n_1$ ,  $\mu_0$ ,  $\mu_1$ ,  $\sigma_0^2$ ,  $\sigma_1^2$ ,  $\pi_0$ ,  $\pi_1$ .

```
n0 <- nrow(data0)
mu0 <- with(data0,mean(x))
var0 <- with(data0,var(x))

n1 <- nrow(data1)
mu1 <- with(data1,mean(x))
var1 <- with(data1,var(x))

n <- n0+n1
pi0 <- n0/n
pi1 <- n1/n</pre>
```

Use  $\sigma_0^2$  and  $\sigma_1^2$  to get an good estimate of the (common) variance.

```
sigma2 \leftarrow 1/(n-2)*((n0-1)*var0+(n1-1)*var1)
```

#### Build your own LDA classifier

Given all these parameters, you can directly define the two linear discriminant functions.

```
discr0 <- function(x){
    x*mu0/sigma2-mu0^2/(2*sigma2)+log(pi0)
}
discr1 <- function(x){
    x*mu1/sigma2-mu1^2/(2*sigma2)+log(pi1)
}</pre>
```

Now build a classifier function, ldaClassifier. This is function of the input variable x and returns a classification based on the values of the discriminant functions.

```
ldaClassifier <- function(x){
  p0 <- discr0(x)
  p1 <- discr1(x)
  return(which.max(c(p0,p1))-1)
}</pre>
```

Test it out on some values...

```
ldaClassifier(-1)
```

```
## [1] 0
```

```
ldaClassifier(2)
```

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

Apply it to the entire data frame.

```
data.df <- data.df %>%
  rowwise() %>%
  mutate(class1 = ldaClassifier(x))
```

How well did it work?

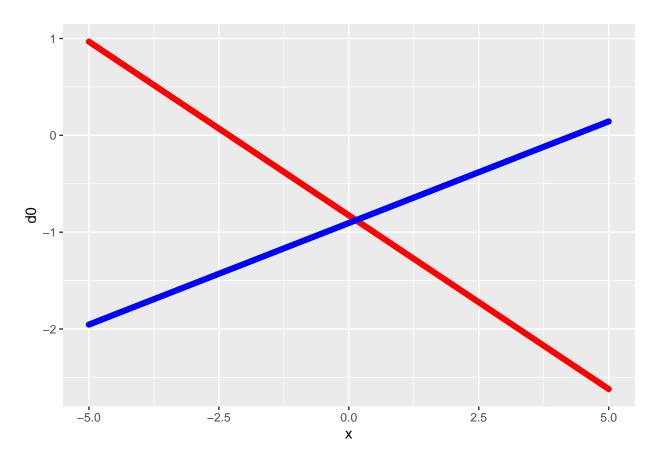
```
with(data.df,table(class,class1))
```

```
## class 0 1
## class 0 1
## 0 44 12
## 1 15 29
```

What is the decision boundary? In other words, what is the value  $x^*$  such that if  $x < x^*$  then classify in one class and if  $x > x^*$  then classify in the other class?

This is just the place where the discriminant functions are equal!

```
xvals <- seq(-5,5,by=.025)
data.frame(x=xvals,d0=discr0(xvals),d1=discr1(xvals)) %>%
    ggplot()+
    geom_point(aes(x,d0),color="red")+
    geom_point(aes(x,d1),color="blue")
```



```
cmp <- discr0(xvals)<discr1(xvals)
max(xvals[cmp])</pre>
```

## [1] 5

# LDA in R

Now apply the lda function in R

```
library(MASS)
mod.lda <- lda(class ~ x, data=data.df)</pre>
```

```
pred.lda <- predict(mod.lda)
data.df$class.lda <- pred.lda$class</pre>
```

If all went well, this should give the exact same prediction as what you had earlier (class1).

```
with(data.df,table(class.lda,class1))
```

```
## class.lda 0 1
## 0 59 0
## 1 0 41
```