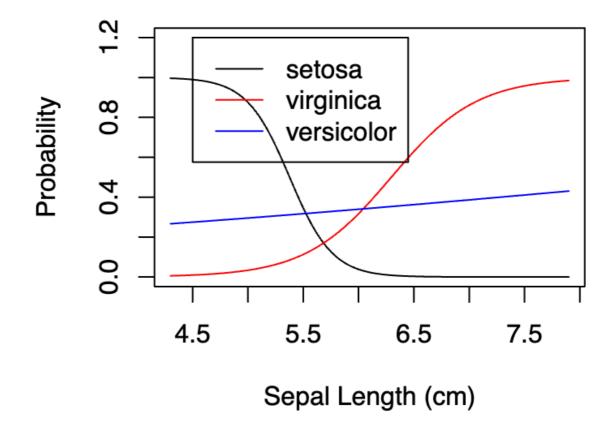
REG Assignment 2

- Sampad Kumar Kar
- MCS202215
- sampadk04@cmi.ac.in

P1



P1.1

If the Sepal.Length` is 6.5 cm, then based on the probability scores as obtained by the Logistic Regression model, virginica is the most likely class, with a probability score of around 0.6`.

P1.2

Yes, Sepal.Length does indeed have an effect on the identification of the class versicolor. Based on the probability plots, we can observe that if the Sepal.Length lies approximately between 5.5 to 6, the model predicts the class versicolor with a probability score of around 0.4.

The linear discriminant rule for class $k \in \{1, 2\}$ is presented as:

$$\delta_k(x) = x^T \Sigma^{-1} \mu_k - rac{1}{2} \mu_k^{-1} \Sigma^{-1} \mu_k + \log(\pi_k)$$

We are also given the following summary statistics from two (independent) datasets:

$$\overline{X}_1=inom{2}{3}$$
 , $\overline{X}_2=inom{5}{7}$ and $S=inom{1}{1}$

Also,
$$S^{-1} = \begin{pmatrix} 2 & -1 \\ -1 & 1 \end{pmatrix}$$
.

Also, $n_1=20$ and $n_2=40$, which means that $\pi_1=\frac{n_1}{n_1+n_2}=\frac{1}{3}$ and $\pi_2=\frac{n_2}{n_1+n_2}=\frac{2}{3}.$

P2.(i)

We compute the linear discriminant function and use that to classify the data point $x=\left(rac{1}{4}
ight)$ as either π_1 or π_2 .

```
In []: # define the function to calculate the delta

lda <- function(x, mu, sigma_inv, pi)
{
    delta <- t(x)%*%sigma_inv%*%mu - (0.5)%*%t(mu)%*%sigma_inv%*%mu + log
    return(delta)
}</pre>
```

```
In []: # setting up the parameters and variables

x <- c(1,4)

mu_1 <- c(2,3)
mu_2 <- c(5,7)

sigma_inv <- matrix(c(2,-1,-1,1), ncol=2, nrow=2, byrow=TRUE)

pi_1 <- 1/3
pi_2 <- 2/3</pre>
```

```
In []: # calculate delta for class 1 and class 2

delta_1 = lda(x, mu_1, sigma_inv, pi_1)
    delta_2 = lda(x, mu_2, sigma_inv, pi_2)

# print out the results
    print(delta_1)
```

print(delta_2)

So, we obtain the following results:

•
$$\delta_1(x) = 1.401388$$

• $\delta_2(x) = -3.905465$

Since, $\delta_1(x) > \delta_2(x)$, we can classify x as π_1 .

P2.(ii)

Under the following assumptions, we can expect our methods to be reliable:

- The data is normally distributed.
- The covariance matrices are equal across all classes.
- The observations are independent of each other.