

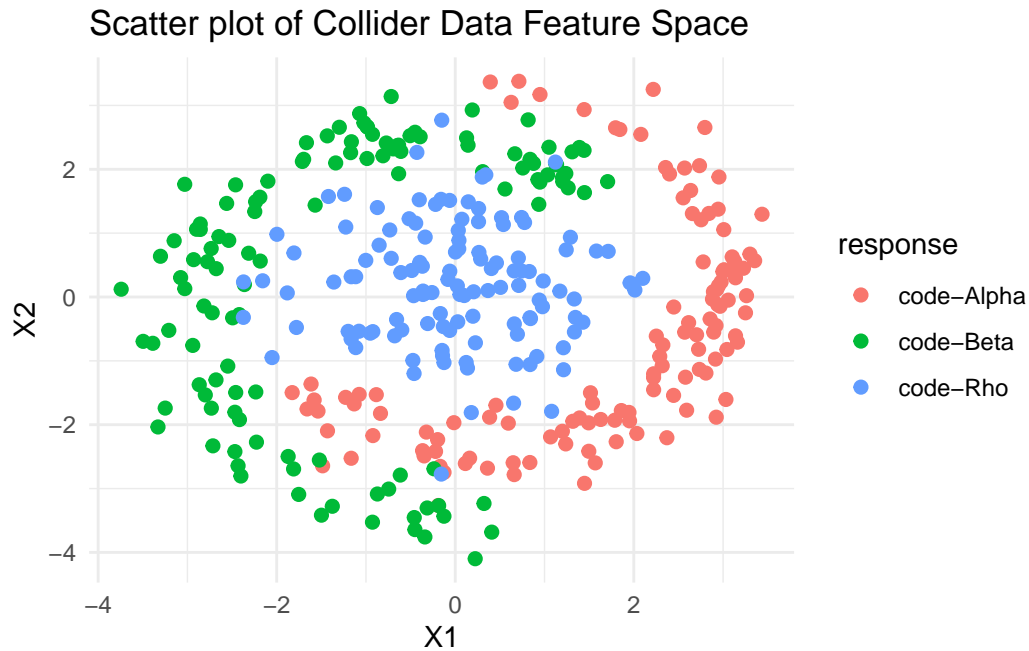
## Assignment 2

```
[1] 360 6
```

	X1	X2	X3	Y1	Y2	Y3
1	3.2509506	-0.2478462	0	1	0	0
2	0.1130716	-2.6077716	1	1	0	0
3	2.0345077	-2.1380898	0	1	0	0
4	-1.6595216	-1.7533739	1	1	0	0
5	3.1474216	0.4139403	1	1	0	0
6	-0.1972298	-2.2335683	0	1	0	0

a)

```
dat$response <- apply(dat[, c("Y1", "Y2", "Y3")], 1, function (x){
  if (x[1] == 1) return("code-Alpha")
  if (x[2] == 1) return("code-Beta")
  if (x[3] == 1) return("code-Rho")
})
ggplot(dat, aes(x=X1, y=X2, color=response ))+geom_point(size=2)+ggtitle("Scatter plot of Co
```



The scatter plot shows how the particles are distributed in the X1-X2 plane. We can see that the process creates a complex and non-linear separation in the points plotted. Therefore, the use of a neural network is justified as they are known for their ability to learn non-linear decision boundaries which is presented in the scatter plot.

**b)**

```
softmax <- function(Z)
{
  Z_shift <- Z - matrix(apply(Z, 2, max),          # subtract column-wise maxima
                        nrow = 3, ncol = ncol(Z),
                        byrow = TRUE)

  expZ    <- exp(Z_shift)                        # exponentiate
  denom   <- matrix(colSums(expZ),                # column-wise sums
                    nrow = 3, ncol = ncol(Z),
                    byrow = TRUE)

  expZ / denom                                  # element-wise division
}
```

c)

```
calc_Ci <- function(y_true, y_pred) {  
  # y_true is assumed to be a scalar (either 0 or 1)  
  if (y_true == 1) {  
    return(-log(y_pred))  
  } else {  
    return(0)  
  }  
}
```

Evaluating only the component corresponding to the actual class simplifies calculations and enhances numerical stability by avoiding evaluating terms that are known to be 0.

d)

```
g <- function(Yhat, Y, eps = 1e-15) {  
  # Yhat, Y : N × q matrices (rows = obs, cols = classes)  
  N <- nrow(Y)  
  -sum( Y * log( pmax(Yhat, eps) ) ) / N  
}
```

e)

number of parameters =  $2p^2 + 2p + 2pm + 2m + m^2 + mq + q$

f)

```
af_forward <- function(X, Y, theta, m, nu)  
{  
  N <- nrow(X)  
  p <- ncol(X)  
  q <- ncol(Y)
```

```

index <- 1:(2*(p^2)) #W1 : p(p+p)
W1 <- matrix(theta[index], nrow=p)

index <- max(index)+1:(2*p) #b1 : (p+p)
b1 <- theta[index]

index <- max(index)+1:((2*p)*m) #W2 : (p+p)*m
W2 <- matrix(theta[index], nrow=2*p)

index <- max(index)+1:m #b2 : m
b2 <- theta[index]

index <- max(index)+1:(m*m) #W3 : (m*m)
W3 <- matrix(theta[index], nrow=m)

index <- max(index)+1:m #b3 : m
b3 <- theta[index]

index <- max(index)+1:(m*q) #W4 : (m*q)
W4 <- matrix(theta[index], nrow=m)

index <- max(index)+1:q #b4 : q
b4 <- theta[index]

#softmax function
softmax <- function(Z)
{
  Z_shift <- Z - matrix(apply(Z, 2, max),          # subtract column-wise maxima
                        nrow = 3, ncol = ncol(Z),
                        byrow = TRUE)

  expZ    <- exp(Z_shift)                          # exponentiate
  denom   <- matrix(colSums(expZ),                 # column-wise sums
                    nrow = 3, ncol = ncol(Z),
                    byrow = TRUE)

  expZ / denom                                     # element-wise division
}

#forward propagation
H1 <- tanh( X %*% W1 + matrix(b1, N, 2*p, TRUE) ) # aug-layer

```

```

H2 <- tanh( H1 %*% W2 + matrix(b2, N, m, TRUE) )      # 2nd hidden
H3 <- tanh(H2 %*% W3 + matrix(b3, N, m, TRUE))
Z <- H3 %*% W4 + matrix(b4, N, q, TRUE)              # logits
probs <- softmax(Z)                                  # softmax

#losses & objective
loss <- g(probs, Y)                                  # cross-entropy
obj <- loss + (nu / 2) * sum(theta^2)
list(probs = probs, loss = loss, obj = obj)
}

```