

Pokemon

```
library(readr)
Pokemon <- read.csv("Pokemon.csv")
View(Pokemon)
names(Pokemon)

## [1] "X."          "Name"        "Type.1"      "Type.2"      "Total"
## [6] "HP"          "Attack"      "Defense"     "Sp..Atk"     "Sp..Def"
## [11] "Speed"       "Generation"  "Legendary"

Pokemon_data <- as.matrix(Pokemon[,c(6:11)])

##Peform PCA
colMeans(Pokemon_data)

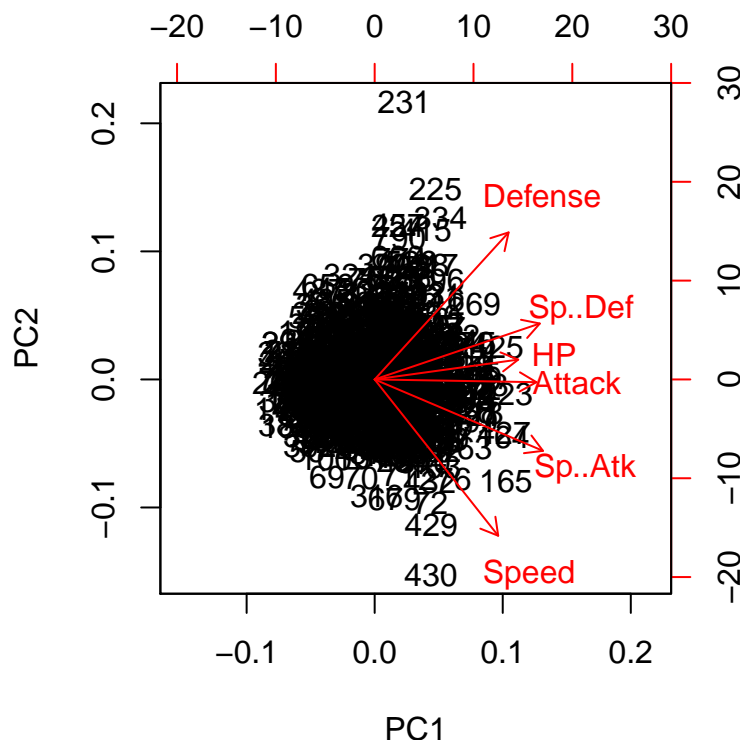
##      HP      Attack      Defense      Sp..Atk      Sp..Def      Speed
## 69.25875 79.00125 73.84250 72.82000 71.90250 68.27750

pok.pr <- prcomp(Pokemon_data,scale=TRUE)

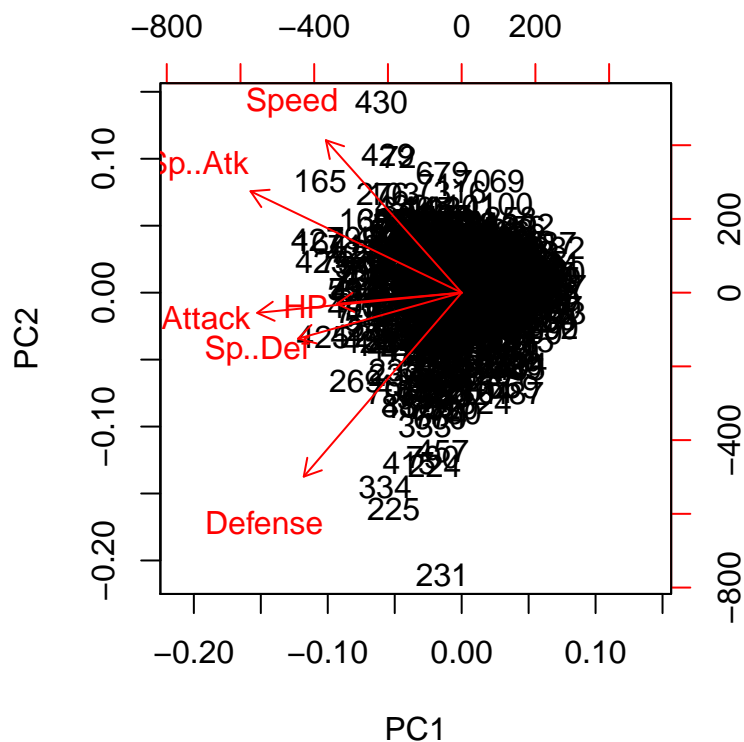
summary(pok.pr)

## Importance of components:
##              PC1      PC2      PC3      PC4      PC5      PC6
## Standard deviation  1.6466  1.0457  0.8825  0.8489  0.65463  0.51681
## Proportion of Variance 0.4519  0.1822  0.1298  0.1201  0.07142  0.04451
## Cumulative Proportion 0.4519  0.6342  0.7640  0.8841  0.95549  1.00000

# Interpreting PCA results with scale=TRUE
biplot(pok.pr)
```



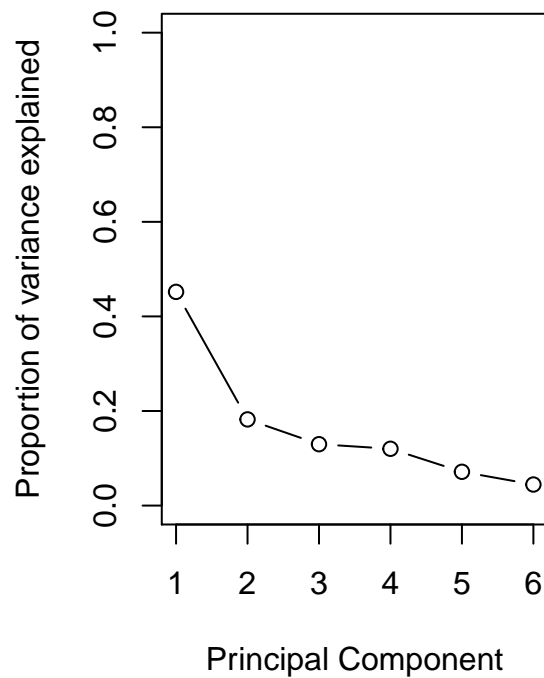
```
#Interpreting PCA results with scale=FALSE
pok.pr.ns <- prcomp(Pokemon_data,scale=FALSE)
biplot(pok.pr.ns)
```



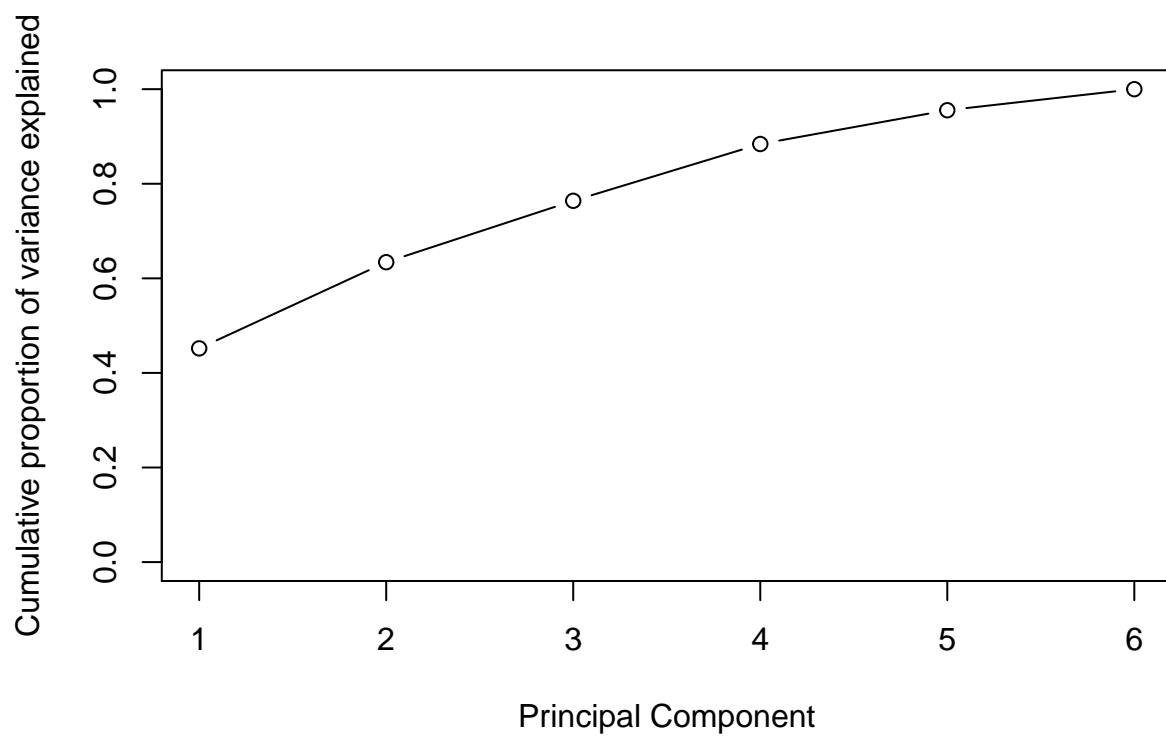
```
##Calculating variability of each component
pr.var <- pok.pr$sdev ^2

#Variance explained by each PC
pve <- pr.var/sum(pr.var)

# Plotting variance explained by each component
par(mfrow=c(1,2))
plot(pve,xlab="Principal Component",ylab="Proportion of variance explained",ylim=c(0,1),type="b")
```



```
plot(cumsum(pve),xlab="Principal Component" ,ylab="Cumulative proportion of variance explained",ylim=c(0,1))
```

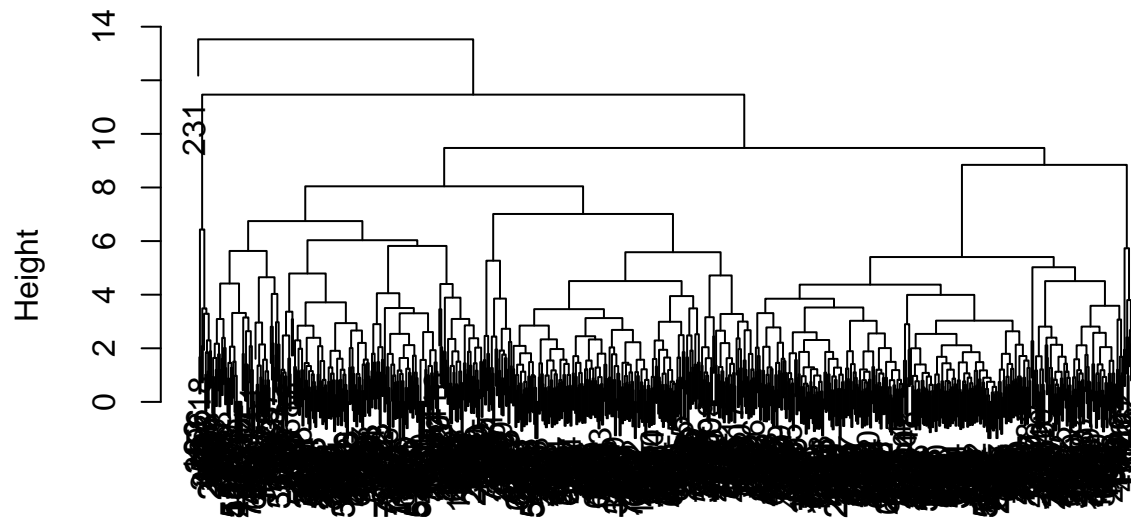


```
#Performing hierarchical clustering using euclidean distance
scaled_data <- scale(Pokemon_data)

Pokemon_hclust<- hclust(dist(scaled_data),method="complete")

plot(Pokemon_hclust)
```

Cluster Dendrogram

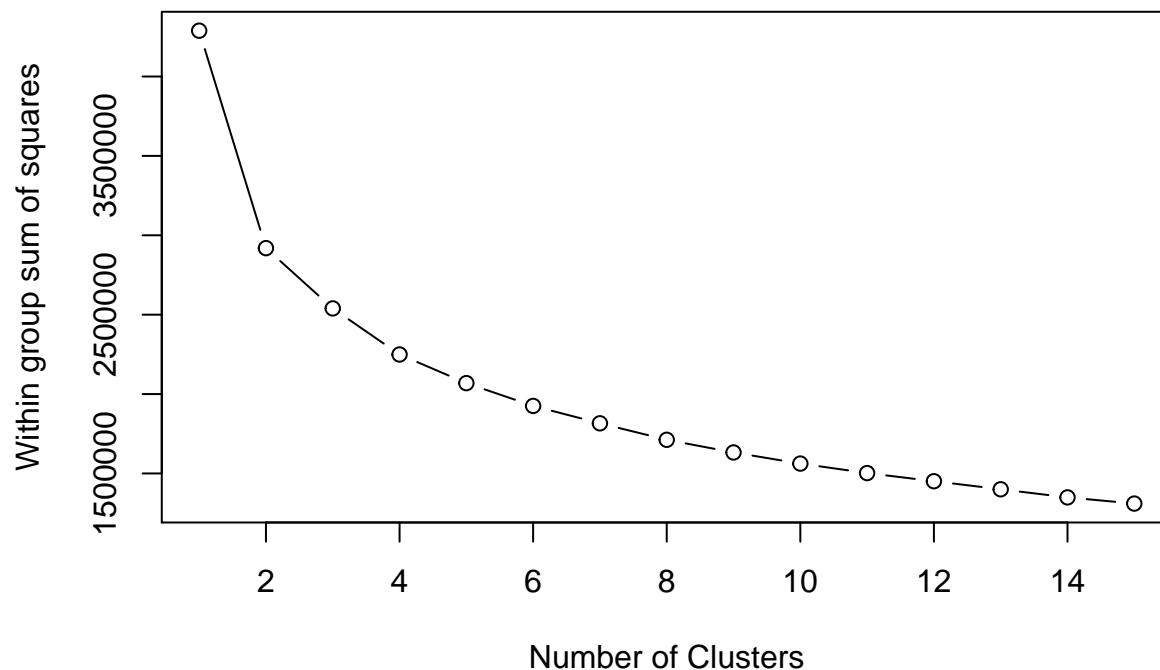


```
dist(scaled_data)
hclust (*, "complete")
```

```
# selecting number of clusters
Pokemon_hclust_clusters <- cutree(Pokemon_hclust,k=4)

#intialize wss
wss <- 0
for(i in 1:15){

  km.out<- kmeans(Pokemon_data,centers = i,nstart = 20,iter.max=50)
  wss[i] <- km.out$tot.withinss
}
plot(1:15,wss,type="b",xlab="Number of Clusters",ylab="Within group sum of squares")
```



```
k <- 4

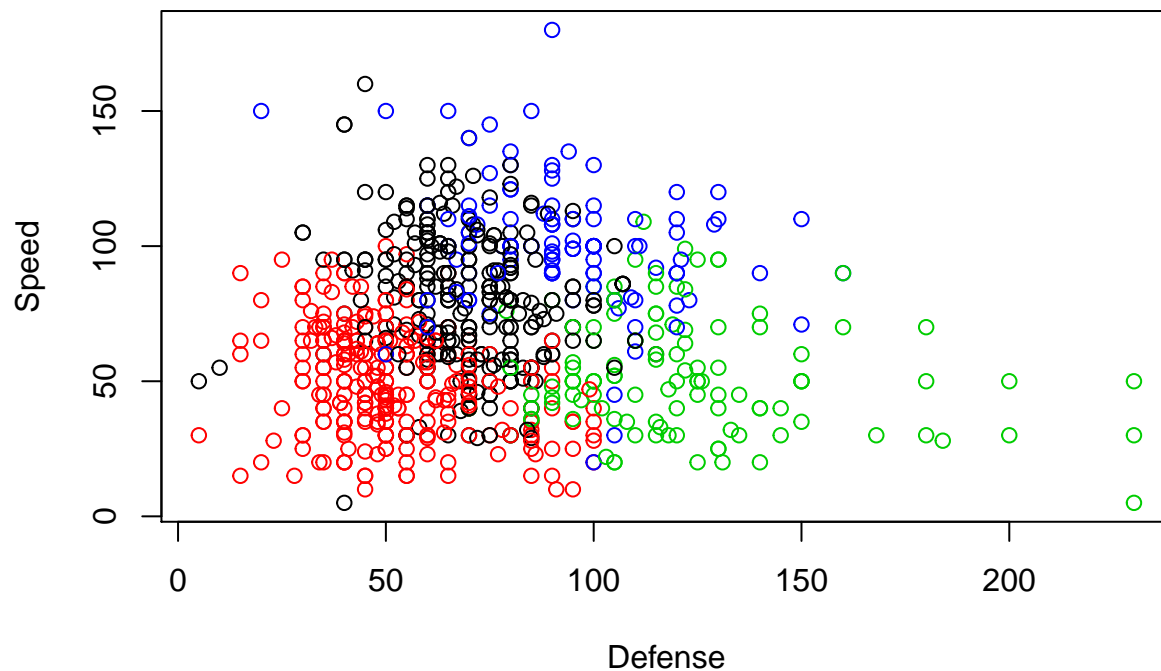
# Build model with k clusters: km.out
km.out <- kmeans(Pokemon_data, centers = k, nstart = 20, iter.max = 50)
km.out

## K-means clustering with 4 clusters of sizes 288, 283, 115, 114
##
## Cluster means:
##      HP      Attack    Defense  Sp..Atk Sp..Def    Speed
## 1 79.18056  81.31944  69.19097  82.01042 77.53125  80.10417
## 2 50.29682  54.03180  51.62898  47.90459 49.15548  49.74912
## 3 71.30435  92.91304 121.42609  63.89565 88.23478  52.36522
## 4 89.20175 121.09649  92.73684 120.45614 97.67544 100.44737
##
## Clustering vector:
##  [1] 2 1 1 4 2 1 1 4 4 2 1 1 4 2 2 1 2 2 1 1 2 2 1 4 2 1 2 1 2 1 2 1 2 3 2
## [36] 2 1 2 2 1 2 1 2 1 2 1 2 1 2 1 1 2 3 2 1 2 1 2 1 2 1 2 1 2 4 2 2 1 2 1
## [71] 1 4 2 1 3 2 1 1 2 1 2 3 3 1 1 2 3 3 2 1 2 2 1 2 1 2 1 2 3 2 1 1 4 3 2
## [106] 1 2 3 2 1 2 1 2 3 1 3 2 2 3 2 3 1 1 1 4 2 1 2 1 2 1 1 1 1 1 3 4 1 2
## [141] 1 4 1 2 2 1 1 4 1 2 3 2 3 1 4 1 4 4 4 2 1 4 4 4 4 4 2 1 1 2 1 1 2 1 1
## [176] 2 1 2 1 2 1 2 2 1 2 1 2 2 2 2 1 2 1 2 2 1 4 3 2 1 3 1 2 2 1 2 2 1 1 2
## [211] 3 1 3 1 1 1 2 1 1 2 3 1 3 3 3 2 1 1 3 3 3 1 3 1 2 1 2 3 2 1 2 2 1 2 1
## [246] 3 2 1 4 1 2 3 1 1 2 2 3 2 2 2 1 1 4 4 3 2 1 4 4 4 4 4 2 1 1 4 2 1 4 4
## [281] 2 1 1 4 2 1 2 1 2 2 1 2 2 2 2 1 2 2 1 2 1 2 1 2 2 1 4 2 1 2 1 2 1 4 2
## [316] 1 2 2 2 1 2 1 2 3 2 2 2 3 2 3 2 3 3 3 2 1 1 2 1 4 1 1 1 1 1 2 1 2 1 4
## [351] 1 1 2 1 4 3 2 1 2 2 2 1 2 1 2 1 4 1 1 1 1 2 1 2 1 2 3 2 3 2 3 2 1 1 3
## [386] 2 1 4 2 3 1 1 1 4 2 2 1 4 2 1 1 2 3 3 3 2 2 3 4 4 2 3 3 4 3 3 3 4 4 4
## [421] 4 4 4 4 4 4 4 4 4 4 4 4 3 4 2 3 3 2 1 1 2 1 1 2 2 1 2 1 2 2 2 2 1 2 1
## [456] 3 3 2 3 3 3 1 2 3 1 2 1 2 1 2 1 1 2 1 2 1 4 1 1 2 1 2 2 1 2 3 2 2 2 1
## [491] 3 2 1 4 4 1 2 4 4 2 3 2 3 2 1 1 2 1 2 2 1 4 1 3 3 3 3 4 4 1 1 3 1 3 1
## [526] 1 1 4 3 3 1 1 1 1 1 1 1 1 3 4 4 4 4 4 4 4 4 3 1 4 4 4 4 4 4 2 1 1 2 1 1
## [561] 2 1 1 2 1 2 2 3 2 1 2 1 2 1 2 1 2 1 2 2 1 2 1 2 3 3 2 1 2 1 1 3 2 3 3
```

```
## [596] 2 1 1 3 1 2 3 1 2 2 1 2 1 2 1 1 2 2 1 2 1 1 1 2 3 2 3 1 2 3 3 3 1 4 2
## [631] 1 2 1 2 1 2 1 1 2 2 1 2 1 2 1 1 2 1 1 2 3 2 1 2 1 1 2 1 2 3 2 3 3 2 1
## [666] 1 2 1 2 2 1 2 1 4 2 1 1 2 1 1 2 1 3 2 3 2 3 3 2 1 2 3 1 3 2 1 4 2 1 4
## [701] 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 2 3 3 2 1 1 2 1 1 2 1 2 2 1 2 2 1
## [736] 2 1 2 2 1 2 1 2 1 1 2 1 1 2 3 4 3 2 1 2 1 2 1 2 3 2 3 2 1 2 1 2 3 2 1
## [771] 1 1 1 3 2 1 4 1 2 1 2 2 2 2 3 3 3 3 2 3 2 1 4 4 4 3 4 4 4 4 4
##
## Within cluster sum of squares by cluster:
## [1] 871531.3 513476.6 455965.7 408271.9
## (between_SS / total_SS = 47.6 %)
##
## Available components:
##
## [1] "cluster"      "centers"      "totss"        "withinss"
## [5] "tot.withinss" "betweenss"    "size"         "iter"
## [9] "ifault"
```

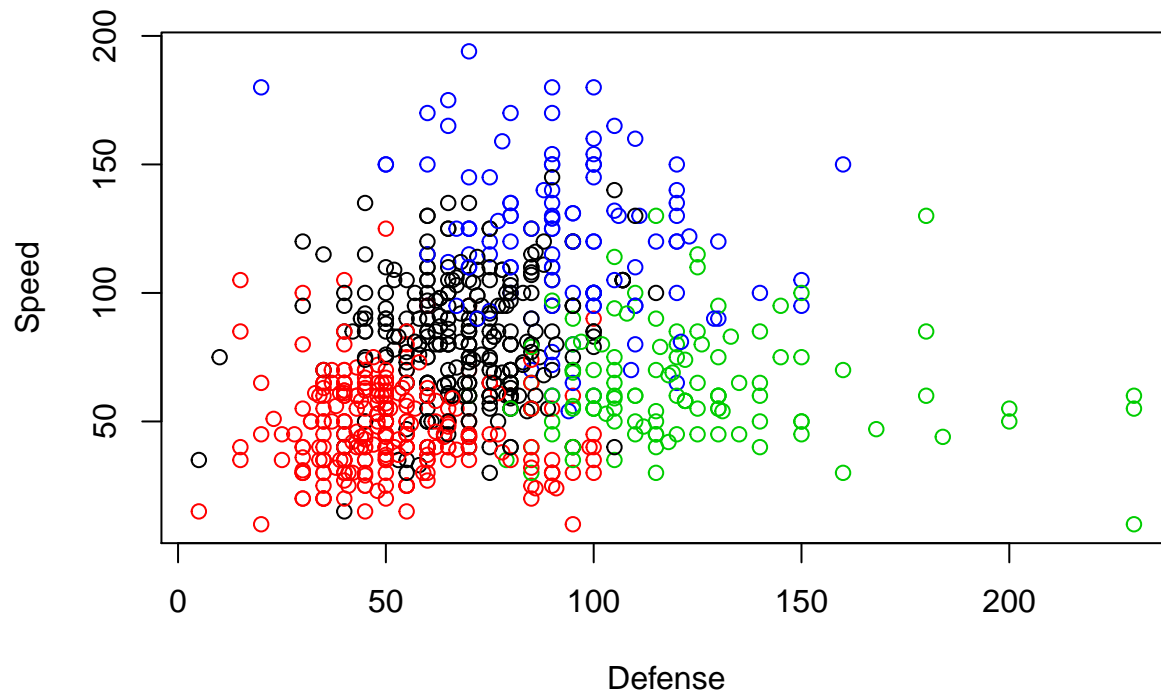
```
# Plot of Defense vs. Speed by cluster membership
plot(Pokemon_data[, c("Defense", "Speed")],
     col = km.out$cluster,
     main = paste("k-means clustering of Pokemon with", k, "clusters"),
     xlab = "Defense", ylab = "Speed")
```

k-means clustering of Pokemon with 4 clusters



```
# Plot of Defense vs. Sp. Atk by cluster membership
plot(Pokemon_data[,3], Pokemon_data[,4],
     col = km.out$cluster,
     main = paste("k-means clustering of Pokemon with", k, "clusters"),
     xlab = "Defense", ylab = "Speed")
```

k-means clustering of Pokemon with 4 clusters



```
table(km.out$cluster,Pokemon_hclust_clusters)
```

```
##      Pokemon_hclust_clusters
##      1  2  3  4
## 1  32 245 11  0
## 2 277  6  0  0
## 3  10 104  0  1
## 4   5 109  0  0
```

```
# Create a hierarchical clustering model: wisc.pr.hclust
pokemon.pr.hclust <- hclust(dist(pok.pr$x[, 1:6]), method = "complete")
```

```
# Cut model into 4 clusters: wisc.pr.hclust.clusters
pokemon.pr.hclust.clusters <- cutree(pokemon.pr.hclust, k = 4)
```