

center;

codigo

September 6, 2020

0.0.1. JUPYTER COSAS

```
[14]: if (!require(cluster)){
      install.packages("cluster", repos='http://cran.us.r-project.org')
    }
    if (!require(purrr)){
      install.packages("purrr", repos='http://cran.us.r-project.org')
    }
    if (!require(fpc)){
      install.packages("fpc", repos='http://cran.us.r-project.org')
    }
    if (!require(dbscan)){
      install.packages("dbscan", repos='http://cran.us.r-project.org')
    }
    if (!require(factoextra)){
      install.packages("factoextra", repos='http://cran.us.r-project.
      .org')
    }
    if (!require(Matrix)){
      install.packages("Matrix", repos='http://cran.us.r-project.org')
    }
    if (!require(xtable)){
      install.packages("xtable", repos='http://cran.us.r-project.org')
    }
    library(cluster)
    library(purrr)
    library(fpc)
    library(dbscan)
    library(factoextra)
    library(Matrix)
    library(xtable)
```

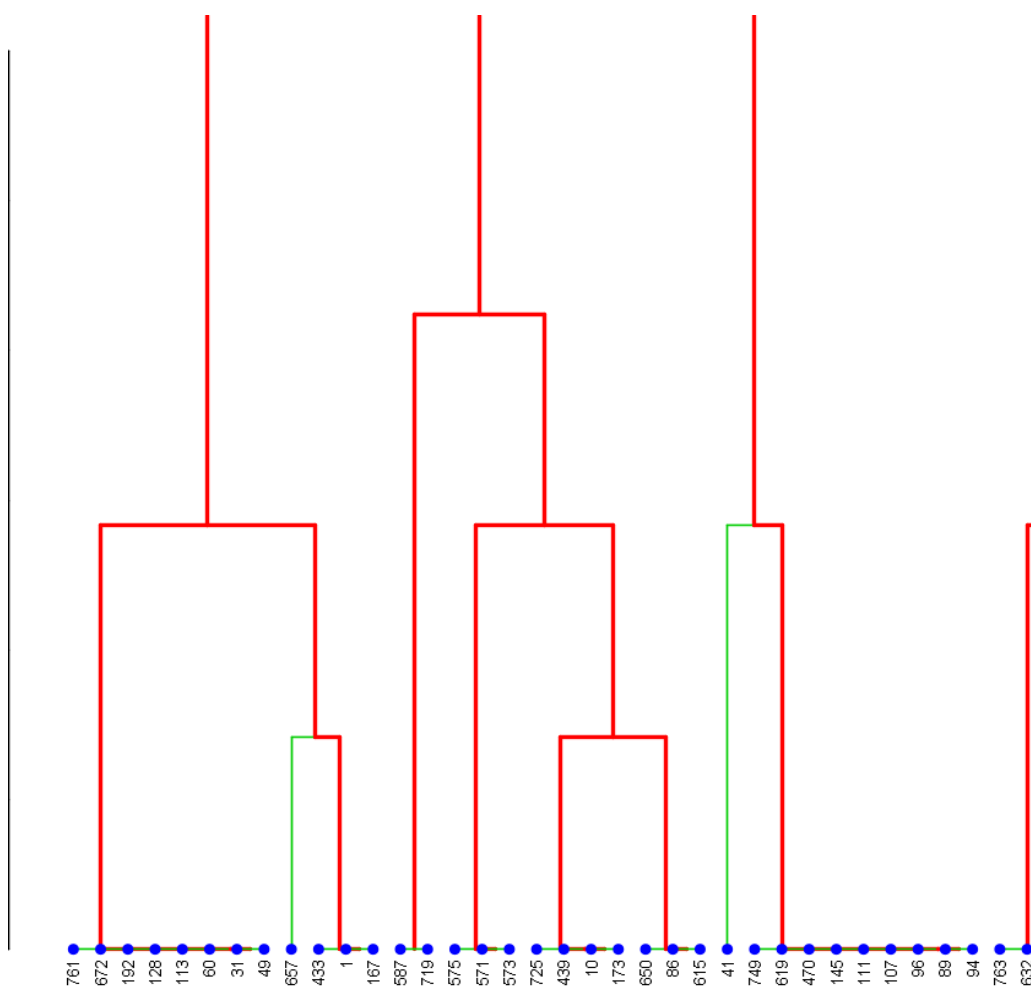
```
[19]: pokemon <- read.csv("Pokemon.csv", encoding="UTF-8", header=TRUE,
      sep=",", strip.white=TRUE)[,c('Name', 'Total')]
    pokemon <- data.frame(pokemon)
    pokemon_fig <- head(pokemon, 10)
    pokemon_fig
```

	Name <fct>	Total <int>
1	Bulbasaur	318
2	Ivysaur	405
3	Venusaur	525
4	VenusaurMega Venusaur	625
5	Charmander	309
6	Charmeleon	405
7	Charizard	534
8	CharizardMega Charizard X	634
9	CharizardMega Charizard Y	634
10	Squirtle	314

A data.frame: 10 × 2

```
[22]: d <- dist(pokemon, method="euclidian")
      hc1 <- hclust(d, method="complete")
      hc1 <- as.dendrogram(hc1)
      par(mar = c(2,0,0,0))
      nodePar <- list(lab.cex=0.6, pch=c(NA, 19), cex=0.7, col="blue")
      plot(hc1, nodePar=nodePar, edgePar=list(col=2:3, lwd=2:1),
            labels=NULL, cex=0.6, xlim=c(0,35), ylim=c(0,6))
```

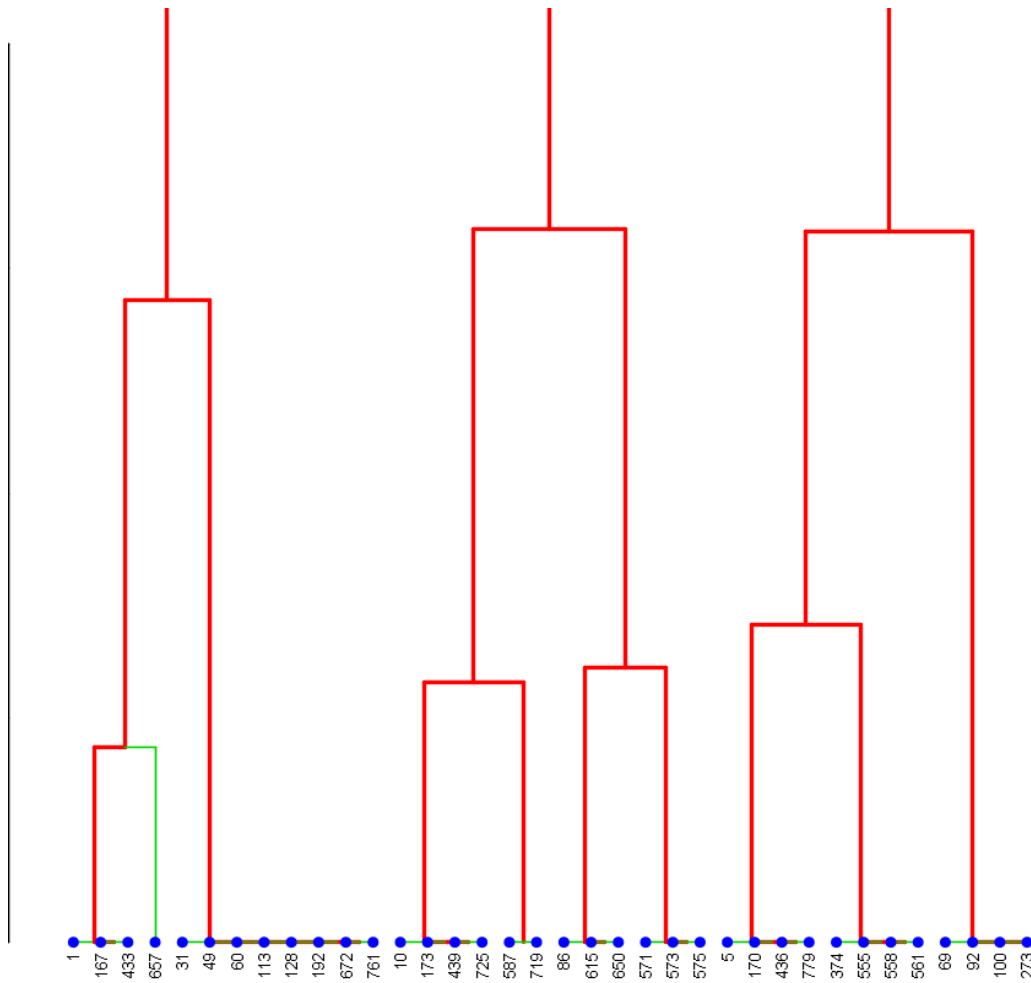
Warning message in dist(pokemon, method = "euclidian"):
 "NAs introducidos por coerción"



```
[24]: hc2 <- agnes(d, method="complete")
m <- c( "average", "single", "complete", "ward")
names(m) <- c( "average", "single", "complete", "ward")
ac <- function(x) {
  agnes(d,method=x)$ac
}
map_dbl(m, ac)
```

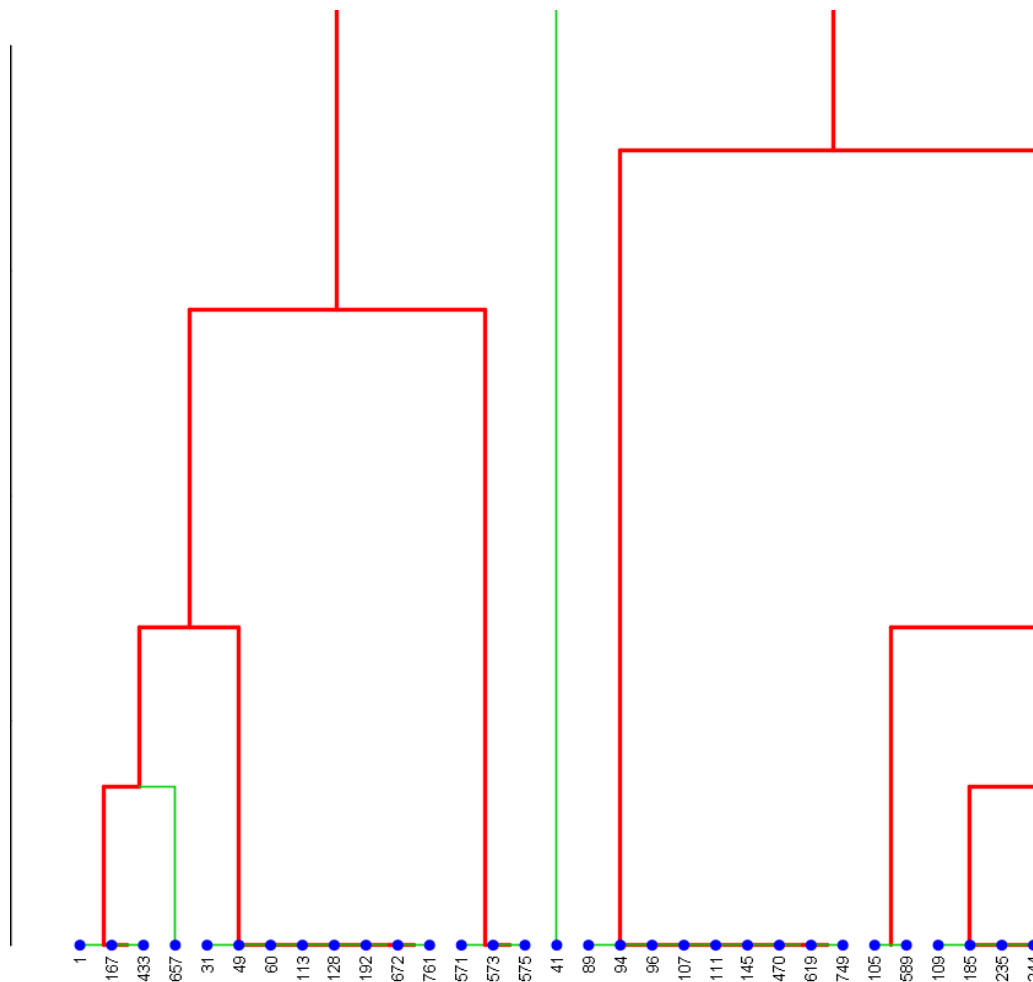
average	0.999318908624144	single	0.996050000000001	complete
	0.999622916666667	ward	0.999933844937563	

```
[27]: hc3 <- as.dendrogram(agnes(d, method="ward"))
      par(mar = c(2,0,0,0))
      plot(hc3, nodePar=nodePar ,edgePar=list(col=2:3, lwd=2:1),
            labels=NULL, cex=0.6, xlim=c(0,35), ylim=c(0,8))
```



```
[28]: hc4 <- diana(d)
      hc4d <- as.dendrogram(hc4)
      par(mar = c(2,0,0,0))
```

```
plot(hc4d, nodePar=nodePar ,edgePar=list(col=2:3, lwd=2:1),  
     labels=NULL, cex=0.6, xlim=c(0,30), ylim=c(0,8))
```



```
[32]: pokemon <- head(pokemon, 100)  
d <- dist(pokemon, method="euclidian")  
hc4 <- diana(d)  
clust <- cutree(hc4, k=5)  
fviz_cluster(list(data = d, cluster=clust))
```

Warning message in dist(pokemon, method = "euclidian"):

"NAs introducidos por coerción"

