

PCA-R

November 19, 2021

```
[ ]: library("tidyverse")
      library("FactoMineR")
      library("factoextra")
      library("FactoInvestigate")
      library("corrplot")

[ ]: df = read_csv('datasets/data.csv')
      head(df)
```

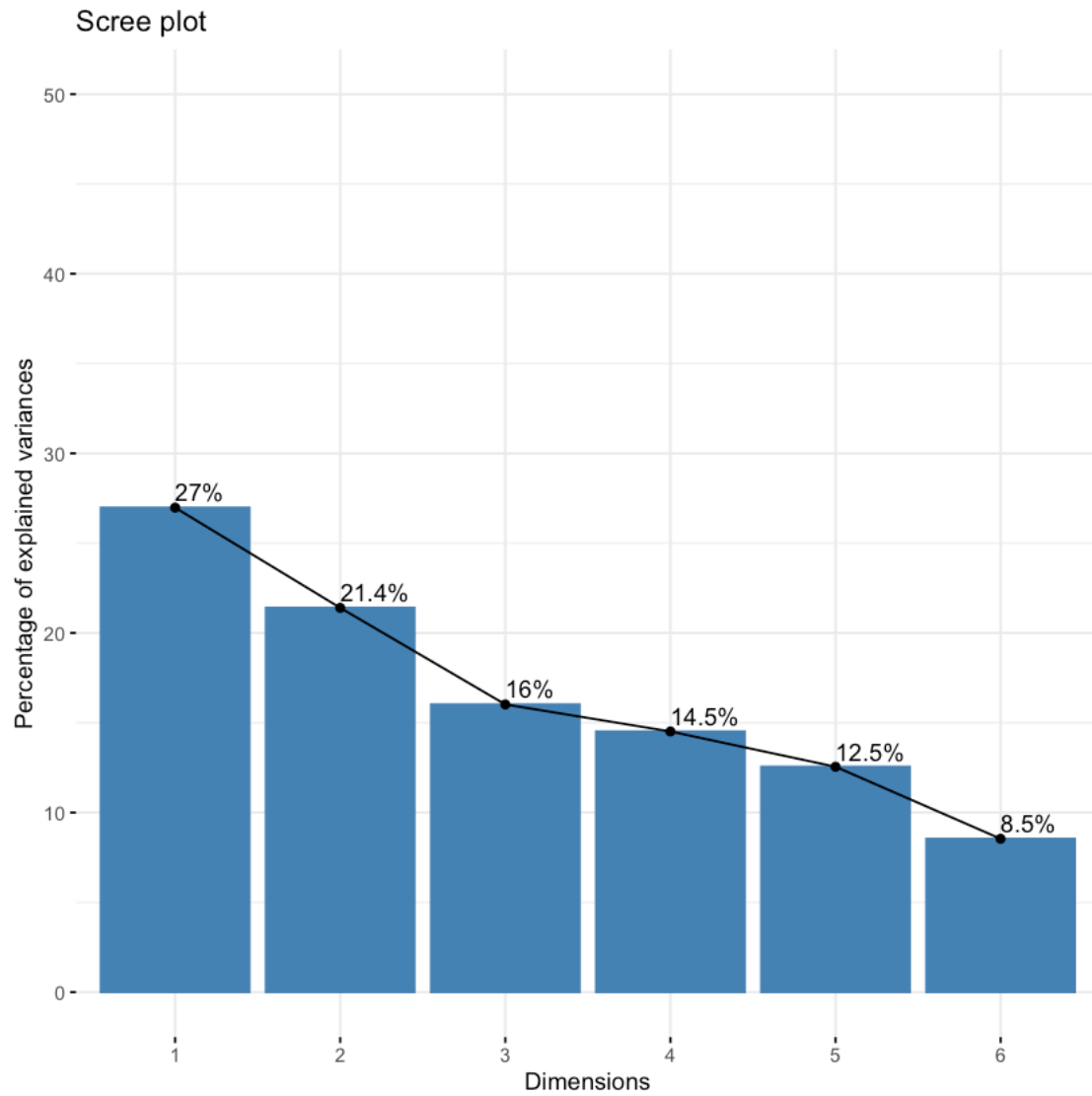
1 PCA

```
[3]: df_pca = df %>% select('score', 'n_matches', 'n_updates_photo', 'n_photos',
      ↪ 'sent_ana', 'length_prof', 'account_age')

      # Apply PCA
      res.pca = PCA(df_pca, scale.unit=TRUE, quanti.sup=1)
```

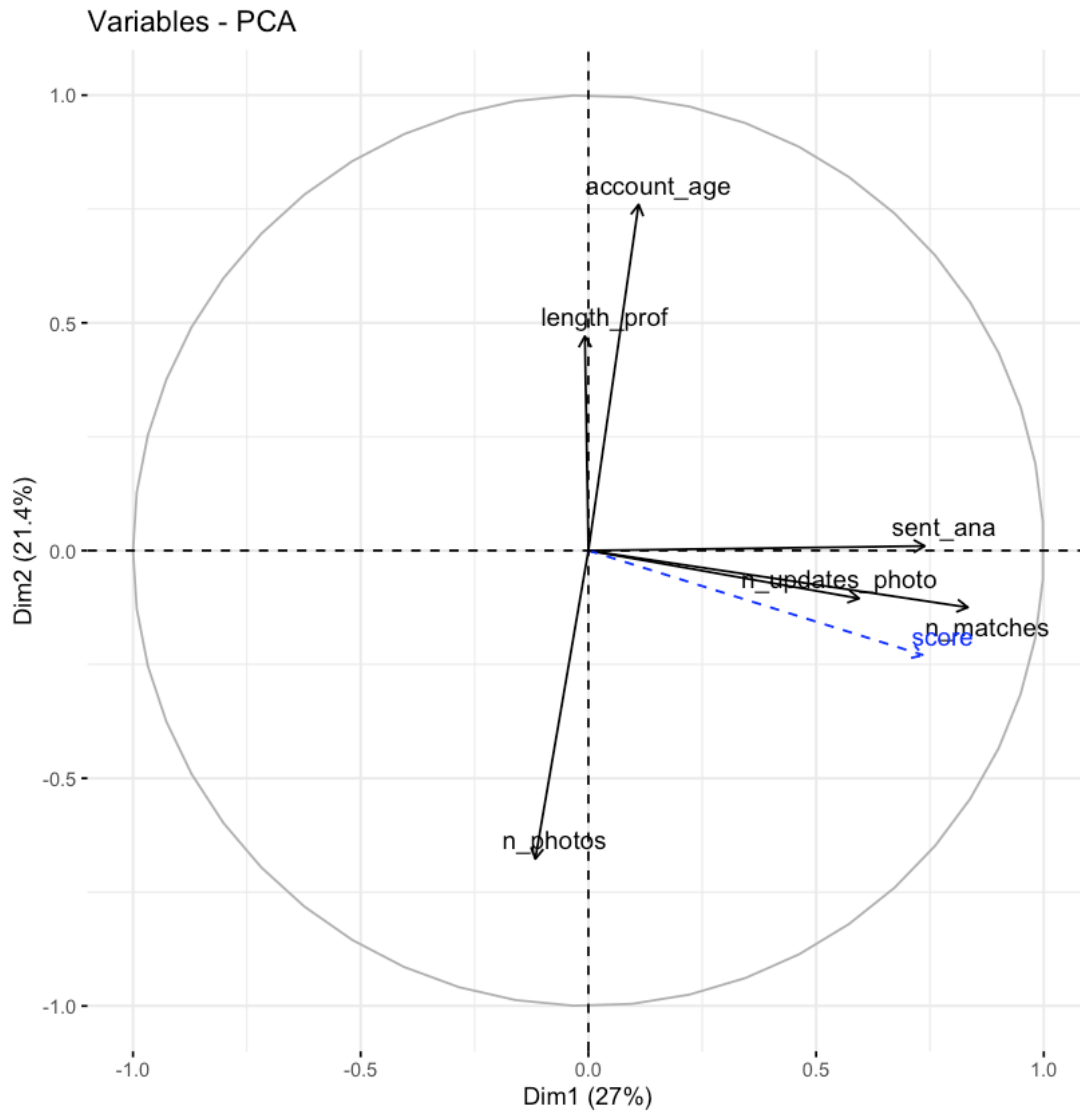
1.0.1 Scree plot

```
[4]: fviz_eig(res.pca, addlabels = TRUE, ylim = c(0, 50), repel = TRUE)
```



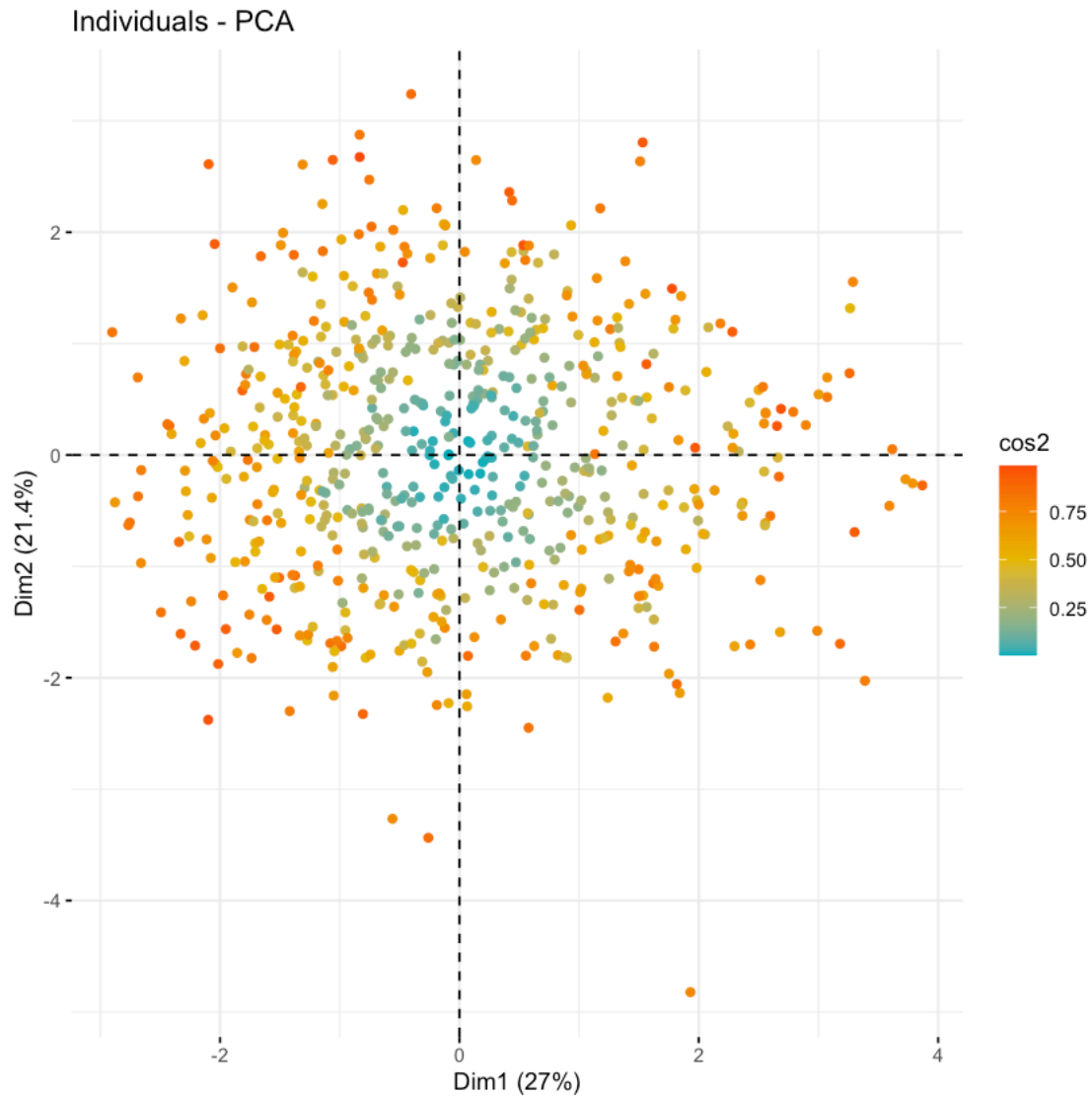
1.0.2 Correlation circle

```
[5]: fviz_pca_var(res.pca, repel = TRUE)
```



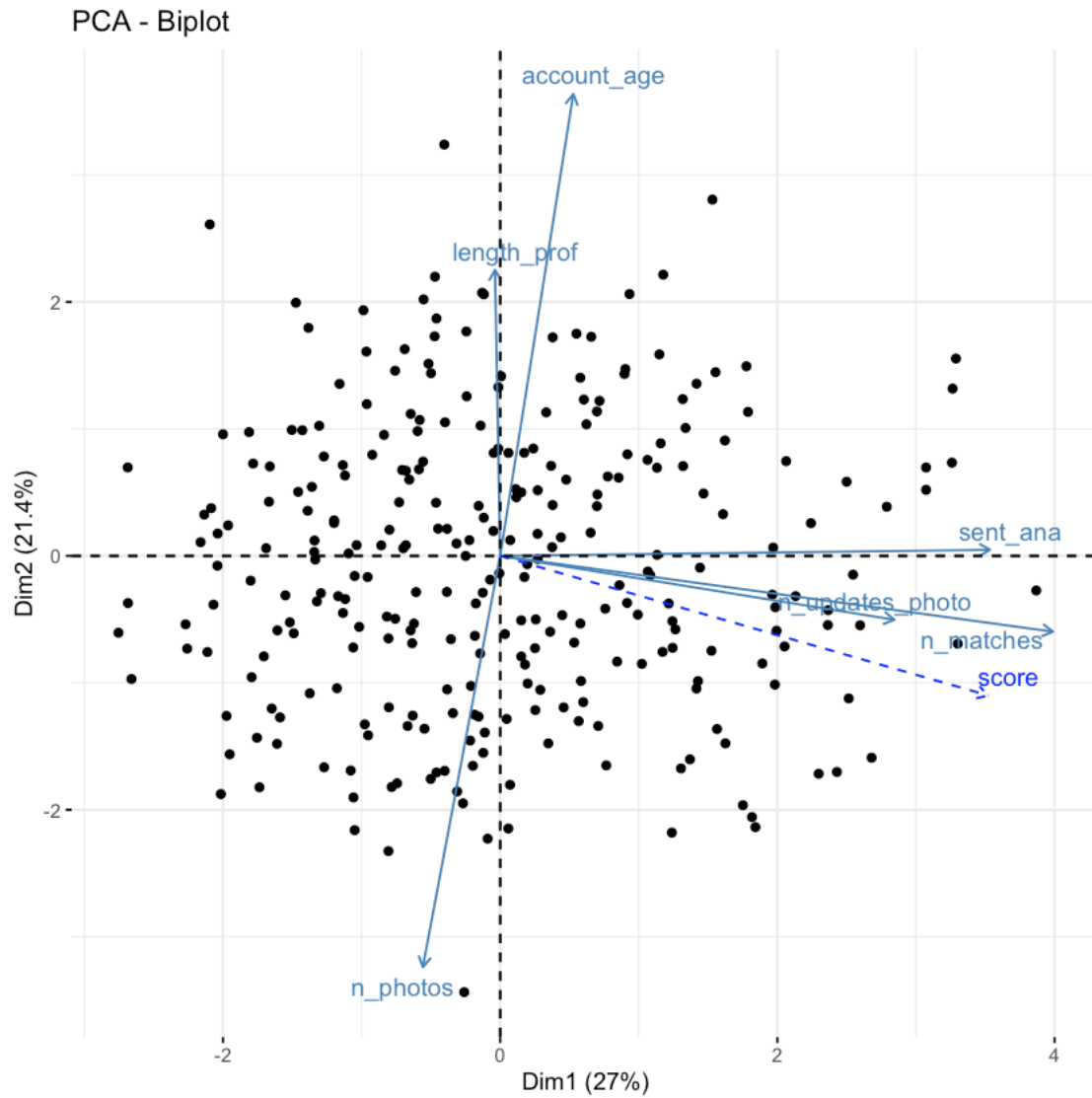
1.0.3 Individual plot

```
[6]: set.seed(123)
# Graph of individuals
fviz_pca_ind(res.pca, col.ind = "cos2",
             gradient.cols = c("#00AFBB", "#E7B800", "#FC4E07"),
             label = "none",
             select.ind = list(name = sample(1:2931, 750, replace=TRUE)), # 25%
             ↪ sample
             )
```



1.0.4 Biplot

```
[7]: set.seed(123)
      # Biplot of individuals and variables
      fviz_pca_biplot(res.pca, label = "var",
                      select.ind = list(name = sample(1:2931, 300, replace=TRUE)), #
      ↪ 10% sample
                      repel = TRUE)
```



1.0.5 Loadings plot of each Principal Component

```
[8]: corrplot(res.pca$var$coord, is.corr=FALSE)
```

