FactoMineR and factoextra: Principal Component Analysis Visualization - R software and data mining

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Principal component analysis (PCA) allows us to summarize the variations (informations) in a data set described by multiple variables. Each variable could be considered as a different dimension. If you have more than 3 variables in your data sets, it could be very difficult to visualize a multi-dimensional hyperspace.

The goal of principal component analysis is to transform the initial variables into a new set of variables which explain the variation in the data. These new variables corresponds to a linear combination of the originals and are called principal components.

PCA reduces the dimensionality of multivariate data, to two or three that can be visualized graphically with minimal loss of information.

Several functions from different packages are available in R for performing PCA: prcomp and princomp (built-in R stats package), PCA (FactoMineR package), dudi.pca(ade4 package).

This R tutorial describes:

1. How to perform a principal component analysis using R software and FactoMineR package

Install and load FactoMineR package

FactoMineR (Husson et al.) is one of the most **powerful R packages** and my favorite one for performing a multivariate exploratory data analysis. A rich documentation is available on the FactoMineR official website (http://factominer.free.fr/index.html) and on youtube. Many thanks to François Husson for this effort...

FactoMineR can be installed and loaded as follow:

```
install.packages("FactoMineR")
library("FactoMineR")
```

Install and load factoextra for visualization

The package <u>factoextra</u> has flexible methods for the classes PCA, prcomp, princomp and dudi in order to extract and visualize quickly the results of the analysis. The ggplot2 plotting system is used for the data visualization.

Install and load **factoextra** as follow:

```
library("devtools")
install_github("kassambara/factoextra")
Load it:
library("factoextra")
```

Prepare the data

We'll use the data sets *decathlon2* from the package **factoextra**:

```
data(decathlon2)
head(decathlon2[, 1:6])
```

	X100m	Long.jump	Shot.put	High.jump	X400m	X110m.hurdle
SEBRLE	11.04	7.58	14.83	2.07	49.81	14.69
CLAY	10.76	7.40	14.26	1.86	49.37	14.05
BERNARD	11.02	7.23	14.25	1.92	48.93	14.99
YURKOV	11.34	7.09	15.19	2.10	50.42	15.31
ZSIVOCZKY	11.13	7.30	13.48	2.01	48.62	14.17
McMULLEN	10.83	7.31	13.76	2.13	49.91	14.38

This data is just a subset of the *decathlon* data in **FactoMineR** package

As illustrated below, the data used here describes athletes' performance during two sporting events (Desctar and OlympicG). It contains 27 individuals (athletes) described by 13 variables:

name	100m	Long.jump	Shot.put	High.jump	400m	110m.hurdk	Discus	Pole.vault	Javeline	1500m	Rank	Points	Competition
SEBRLE	11.04	7.58	14.83	2.07	49.81	14.69	43.75	5.02	63.19	291.7	1	8217	Decastar
CLAY	10.76	7.4	14.26	1.86	49.37	14.05	50.72	4.92	60.15	301.5	2	8122	Decastar
BERNARD	11.02	7.23	14.25	1.92	48.93	14.99	40.87	5.32	62.77	280.1	4	8067	Decastar
rurkov	11.34	7.09	15.19	2.1	50.42	15.31	46.26	4.72	63.44	276.4	5	8036	Decastar
ESIVOCZKY	11.13	7.3	13.48	2.01	48.62	14.17	45.67	4.42	55.37	268	7	8004	Decastar
McMULLEN	10.83	7.31	13.76	2.13	49.91	14.38	44.41	4.42	56.37	285.1	8	7995	Decastar
MARTINEAU	11.64	6.81	14.57	1.95	50.14	14.93	47.6	4.92	52.33	262.1	9	7802	Decastar
HERNU	11.37	7.56	14.41	1.86	51.1	15.06	44.99	4.82	57.19	285.1	10	7733	Decastar
BARRAS	11.33	6.97	14.09	1.95	49.48	14.48	42.1	4.72	55.4	282	11	7708	Decastar
NOOL	11.33	7.27	12.68	1.98	49.2	15.29	37.92	4.62	57.44	266.6	12	7651	Decastar
BOURGUIGN	11.36	6.8	13.46	1.86	51.16	15.67	40.49	5.02	\$4.68	291.7	13	7313	Decastar
iebrie	10.85	7.84	16.36	2.12	48.36	14.05	48.72	5	70.52	280.01	1	8893	OlympicG
Clay	10.44	7.96	15.23	2.06	49.19	14.13	50.11	4.9	69.71	282	2	8820	OlympicG
Carpov	10.5	7.81	15.93	2.09	46.81	13.97	51.65	4.6	55.54	278.11	3	8725	OlympicG
Macey	10.89	7.47	15.73	2.15	48.97	14.56	48.34	4.4	58.46	265.42	4	8414	OlympicG
Warners	10.62	7.74	14.48	1.97	47.97	14.01	43.73	4.9	55.39	278.05	5	8343	OlympicG
Esivoczky	10.91	7.14	15.31	2.12	49.4	14.95	45.62	4.7	63.45	269.54	6	8287	OlympicG
Hernu	10.97	7.19	14.65	2.03	48.73	14.25	44.72	4.8	57.76	264.35	7	8237	OlympicG
Bernard	10.69	7.48	14.8	2.12	49.13	14.17	44.75	4.4	55.27	276.31	9	8225	OlympicG
schwarzl	10.98	7.49	14.01	1.94	49.76	14.25	42.43	5.1	56.32	273.56	10	8102	OlympicG
Pogorelov	10.95	7.31	15.1	2.06	50.79	14.21	44.6	5	53.45	287.63	11	8084	OlympicG
schoenbeck	10.9	7.3	14.77	1.88	50.3	14.34	44,41	5	60.89	278.82	12	8077	OlympicG
Barras	11.14	6.99	14.91	1.94	49.41	14.37	44.83	4.6	64.55	267.09	13	8067	OlympicG
CARPOV	11.02	7.3	14.77	2.04	48.37	14.09	48.95	4.92	50.31	300.2	3	8099	Decastar
WARNERS	11.11	7.6	14.31	1.98	48.68	14.23	41.1	4.92	51.77	278.1	6	8030	Decastar
Vool	10.8	7.53	14.26	1.88	48.81	14.8	42.05	5.4	61.33	276.33	8	8235	OlympicG
Drews	10.87	7.38	13.07	1.88	48.51	14.01	40.11	5	51.53	274.21	19	7926	OlympicG
	Active ind												
	Active var	lables											
	Suppleme	ntary quantiti	ative variab	les									
	Suppleme	ntary qualitat	lve variable										
	Supplementary individuals												

Only some of these individuals and variables will be used to perform the principal component analysis (PCA).

The coordinates of the remaining individuals and variables on the factor map will be **predicted** after the PCA.

In PCA terminology, our data contains:

- Active individuals (in blue, rows 1:23): Individuals that are used during the principal component analysis.
- **Supplementary individuals** (in green, rows 24:27): The coordinates of these individuals will be predicted using the PCA informations and parameters obtained with active individuals/variables
- Active variables (in pink, columns 1:10): Variables that are used for the principal component analysis.
- **Supplementary variables**: As supplementary individuals, the coordinates of these variables will be predicted also.
- **Supplementary continuous variables**: Columns 11 and 12 corresponding respectively to the rank and the points of athletes.
- Supplementary qualitative variables: Column 13 corresponding to the two athlete-tic meetings (2004 Olympic Game or 2004 Decastar). This factor variables will be used to color individuals by groups.

Extract only active individuals and variables for principal component analysis:

decathlon2.active <- decathlon2[1:23, 1:10]
head(decathlon2.active[, 1:6])</pre>

	X100m	Long.jump	Shot.put	High.jump	X400m	X110m.hurdle
SEBRLE	11.04	7.58	14.83	2.07	49.81	14.69
CLAY	10.76	7.40	14.26	1.86	49.37	14.05
BERNARD	11.02	7.23	14.25	1.92	48.93	14.99
YURKOV	11.34	7.09	15.19	2.10	50.42	15.31

Exploratory data analysis

Before principal component analysis, we can perform some exploratory data analysis such as descriptive statistics, correlation matrix and scatter plot matrix.

Descriptive statistics

```
decathlon2.active_stats <- data.frame(</pre>
  Min = apply(decathlon2.active, 2, min), # minimum
  Q1 = apply(decathlon2.active, 2, quantile, 1/4), # First quartile
 Med = apply(decathlon2.active, 2, median), # median
 Mean = apply(decathlon2.active, 2, mean), # mean
  Q3 = apply(decathlon2.active, 2, quantile, 3/4), # Third quartile
 Max = apply(decathlon2.active, 2, max) # Maximum
decathlon2.active_stats <- round(decathlon2.active_stats, 1)</pre>
head(decathlon2.active_stats)
                    Q1 Med Mean
                                   Q3 Max
X100m
             10.4 10.8 11.0 11.0 11.2 11.6
Long.jump
              6.8 7.2 7.3 7.3 7.5 8.0
Shot.put
             12.7 14.2 14.7 14.6 15.1 16.4
High.jump
             1.9 1.9 2.0 2.0 2.1 2.1
             46.8 49.0 49.4 49.4 50.0 51.2
X110m.hurdle 14.0 14.2 14.4 14.5 14.9 15.7
```

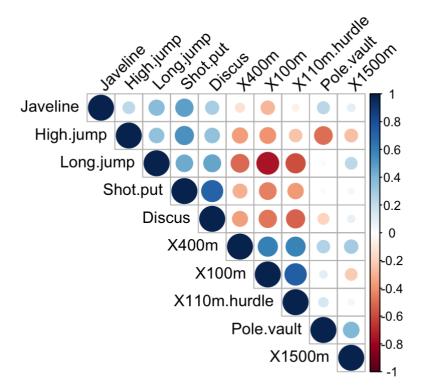
Note that, you can also use the built-in R function **summary()** for the descriptive statistics but I don't like the format of the output on data frame.

Correlation matrix

The correlation between variables can be calculated as follow:

```
cor.mat <- round(cor(decathlon2.active),2)</pre>
head(cor.mat[, 1:6])
             X100m Long.jump Shot.put High.jump X400m X110m.hurdle
                               -0.45
                                          -0.40 0.59
X100m
              1.00
                       -0.76
             -0.76
                        1.00
                                 0.44
                                           0.34 -0.51
                                                              -0.59
Long.jump
             -0.45
                        0.44
                                           0.53 -0.31
Shot.put
                                 1.00
                                                              -0.38
High.jump
             -0.40
                       0.34
                                 0.53
                                           1.00 -0.37
                                                              -0.25
                                           -0.37 1.00
              0.59
                                                               0.58
X400m
                       -0.51
                                -0.31
                                           -0.25 0.58
X110m.hurdle 0.73
                       -0.59
                                -0.38
                                                               1.00
```

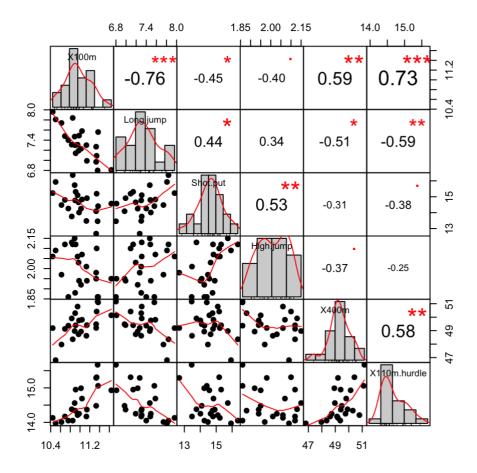
Visualize the correlation matrix using a correlogram: the package corrplot is required.



Read more about visualizing correlation matrix: <u>Correlation matrix visualization</u>

Make a scatter plot matrix showing the correlation coefficients between variables and the significance levels: the package PerformanceAnalytics is required.

```
# install.packages("PerformanceAnalytics")
library("PerformanceAnalytics")
chart.Correlation(decathlon2.active[, 1:6], histogram=TRUE, pch=19)
```



You can read more about this plot here: Correlation matrix visualization

Principal component analysis

The function **PCA()** [in *FactoMiner* package] can be used. A simplified format is :

```
PCA(X, scale.unit = TRUE, ncp = 5, graph = TRUE)
```

- X: a data frame. Rows are individuals and columns are numeric variables
- **scale.unit**: a logical value. If *TRUE*, the data are scaled to unit variance before the analysis. This standardization to the same scale avoids some variables to become dominant just because of their large measurement units.
- ncp: number of dimensions kept in the final results.
- graph: a logical value. If TRUE a graph is displayed.

In the R code below, the PCA is performed only on the active individuals/variables:

```
library("FactoMineR")
res.pca <- PCA(decathlon2.active, graph = FALSE)</pre>
```

The output of the function **PCA()** is a list including:

```
rrint(res.pca)

**Results for the Principal Component Analysis (PCA)**
The analysis was performed on 23 individuals, described by 10 variables
```

*The results are available in the following objects:

```
description
   name
   "$eig"
                       "eigenvalues"
1
  "$var"
                       "results for the variables"
2
3
  "$var$coord"
                      "coord. for the variables"
                      "correlations variables - dimensions"
  "$var$cor"
5
   "$var$cos2"
                      "cos2 for the variables"
  "$var$cos∠
"$var$contrib"
                      "contributions of the variables"
6
   "$ind"
                      "results for the individuals"
7
  "$ind$coord"
"$ind$cos2"
                      "coord. for the individuals"
8
                      "cos2 for the individuals"
9 "$1nd$cosz
10 "$ind$contrib"
                      "contributions of the individuals"
11 "$call"
                      "summary statistics"
12 "$call$centre"
                      "mean of the variables"
13 "$call$ecart.type" "standard error of the variables"
14 "$call$row.w"
                       "weights for the individuals"
15 "$call$col.w"
                       "weights for the variables"
```

The object that is created using the function **PCA()** contains many informations found in many different lists and matrices. These values are described in the next section.

Variances of the principal components

The proportion of variances retained by the principal components can be extracted as follow:

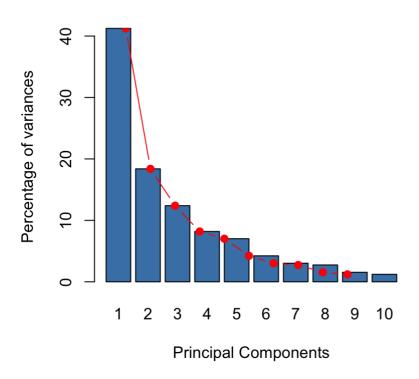
```
eigenvalues <- res.pca$eig
head(eigenvalues[, 1:2])

eigenvalue percentage of variance
comp 1 4.1242133 41.242133
comp 2 1.8385309 18.385309
comp 3 1.2391403 12.391403
comp 4 0.8194402 8.194402
comp 5 0.7015528 7.015528
comp 6 0.4228828 4.228828
```

- **Eigenvalues** correspond to the amount of the variation explained by each principal component (PC). Eigenvalues are large for the first PC and small for the subsequent PCs.
- A PC with an eigenvalue > 1 indicates that the PC accounts for more variance than accounted by one of the original variables in standardized data. This is commonly used as a cutoff point to determine the number of PCs to retain.

Make a scree plot using base graphics: A scree plot is a graph of the eigenvalues/variances associated with components.

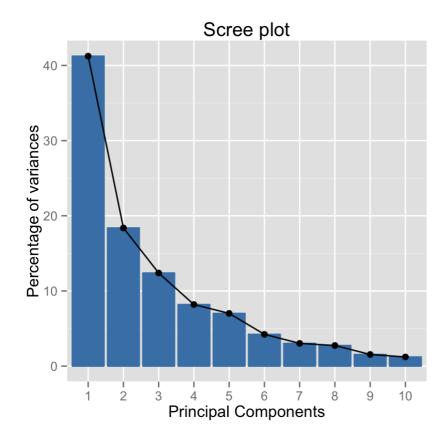
Variances



 \sim 60% of the informations (variances) contained in the data are retained by the first two principal components.

Make the scree plot using the package factoextra:

fviz_screeplot(res.pca, ncp=10)



Graph of individus and variables

The function **plot.PCA()** can be used. A simplified format is:

```
plot.PCA(x, axes = c(1,2), choix=c("ind", "var"))
```

- x : An object of class PCA
- axes: A numeric vector of length 2 specifying the component to plot
- **choix**: The graph to be plotted. Possible values are "ind" for the individuals and "var" for the variables

Variables factor map: The correlation circle

Coordinates of variables on the principal components

head(res.pca\$var\$coord)

	Dim.1	Dim.2	Dim.3	Dim.4	Dim.5
X100m	-0.8506257	-0.17939806	0.3015564	0.03357320	-0.1944440
Long.jump	0.7941806	0.28085695	-0.1905465	-0.11538956	0.2331567
Shot.put	0.7339127	0.08540412	0.5175978	0.12846837	-0.2488129
High.jump	0.6100840	-0.46521415	0.3300852	0.14455012	0.4027002
X400m	-0.7016034	0.29017826	0.2835329	0.43082552	0.1039085
X110m.hurdle	-0.7641252	-0.02474081	0.4488873	-0.01689589	0.2242200

Cos2: quality of variables on the factor map

The quality of representation of the variables of the principal components are called the cos2.

head(res.pca\$var\$cos2)

```
Dim.1Dim.2Dim.3Dim.4Dim.5X100m0.72356410.03218366410.090936280.00112715970.03780845Long.jump0.63072290.07888062850.036307980.01331475060.05436203Shot.put0.53862790.00729386360.267907490.01650412110.06190783High.jump0.37220250.21642420700.108956220.02089473750.16216747X400m0.49224730.08420342090.080390910.18561062690.01079698X110m.hurdle0.58388730.00061210770.201499840.00028547120.05027463
```

Contributions of the variables to the principal components

Variable contributions in the determination of a given principal component are (in percentage): (var.cos2 * 100) / (total cos2 of the component)

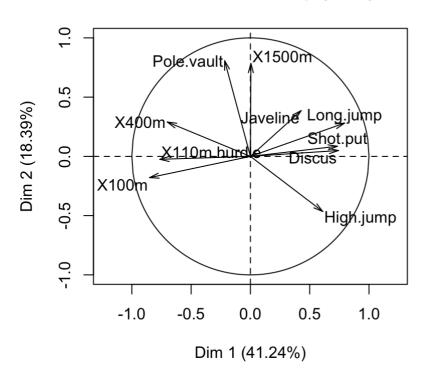
head(res.pca\$var\$contrib)

```
Dim.1Dim.2Dim.3Dim.4Dim.5X100m17.5442931.75050987.3386590.137552405.389252Long.jump15.2931684.29041622.9300941.624859367.748815Shot.put13.0601370.396722421.6204322.014072698.824401High.jump9.02481111.77158388.7928882.5498795123.115504X400m11.9355444.57992966.48763622.650905991.539012X110m.hurdle14.1575440.033293316.2612610.034837357.166193
```

Graph of variables using FactoMineR base graph

plot(res.pca, choix = "var")

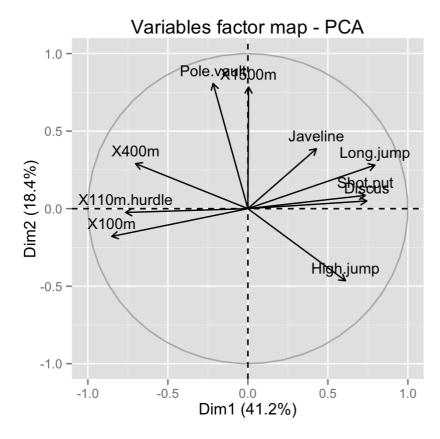
Variables factor map (PCA)



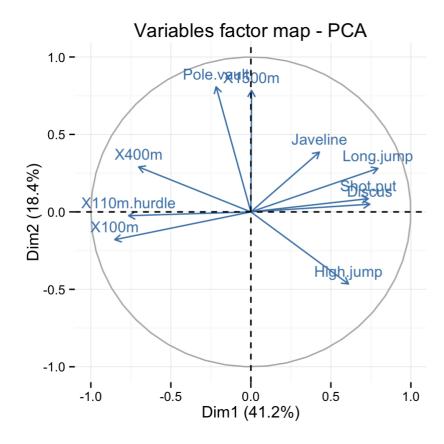
Graph of variables using factoextra

The function fviz_pca_var() is used to visualize variables :

```
# Default plot
fviz_pca_var(res.pca)
```

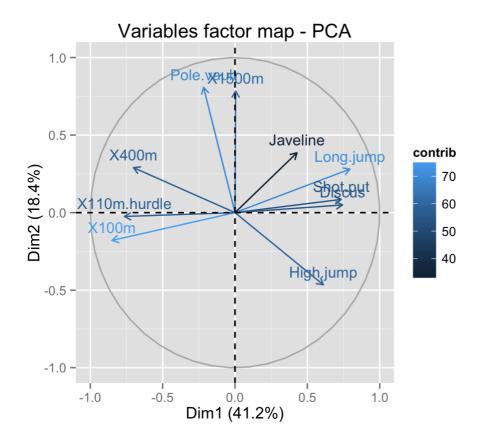


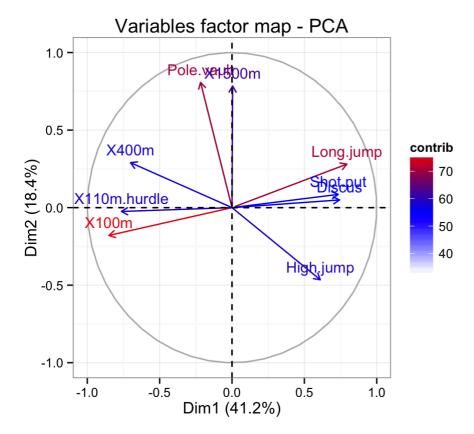
Change color and theme
fviz_pca_var(res.pca, col.var="steelblue")+
 theme_minimal()



Note that, using **factoextra** package, the color or the transparency of variables can be automatically controlled by the value of their contributions, their cos2, their coordinates on x or y axis.

```
# Control variable colors using their contribution
# Possible values for the argument col.var are :
    # "cos2", "contrib", "coord", "x", "y"
fviz_pca_var(res.pca, col.var="contrib")
```

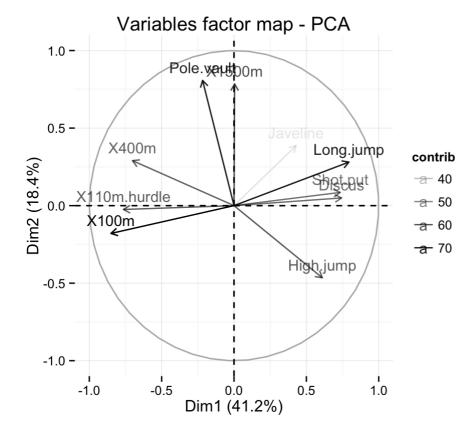




This is helpful to highlight the most important variables in the determination of the principal components.

It's also possible to control automatically the transparency of variables by their contributions:

```
# Control the transparency of variables using their contribution
# Possible values for the argument alpha.var are :
    # "cos2", "contrib", "coord", "x", "y"
fviz_pca_var(res.pca, alpha.var="contrib")+
    theme_minimal()
```



Read more about ggplot2 and colors here : <u>ggplot2 colors - How to change colors automatically and manually?</u>

Graph of individuals

Coordinates of individuals on the principal components

head(res.pca\$ind\$coord)

	Dim.1	Dim.2	Dim.3	Dim.4	Dim.5
SEBRLE	0.1955047	1.5890567	0.6424912	0.08389652	1.16829387
CLAY	0.8078795	2.4748137	-1.3873827	1.29838232	-0.82498206
BERNARD	-1.3591340	1.6480950	0.2005584	-1.96409420	0.08419345
YURKOV	-0.8889532	-0.4426067	2.5295843	0.71290837	0.40782264
ZSIVOCZKY	-0.1081216	-2.0688377	-1.3342591	-0.10152796	-0.20145217
McMULLEN	0.1212195	-1.0139102	-0.8625170	1.34164291	1.62151286

Cos2: quality of representation of individuals on the principal components

head(res.pca\$ind\$cos2)

```
Dim.1Dim.2Dim.3Dim.4Dim.5SEBRLE0.0075301790.497473230.0813252320.0013866880.2689026575CLAY0.0487012490.457016600.1436281170.1257917410.0507850580BERNARD0.1971998040.289965550.0042940150.4118191830.0007567259YURKOV0.0961098000.023825710.7782303220.0618126370.0202279796ZSIVOCZKY0.0015743850.576419440.2397541520.0013882160.0054654972McMULLEN0.0021754370.152194990.1101378720.2664865300.3892621478
```

Contribition of individuals to the princial components

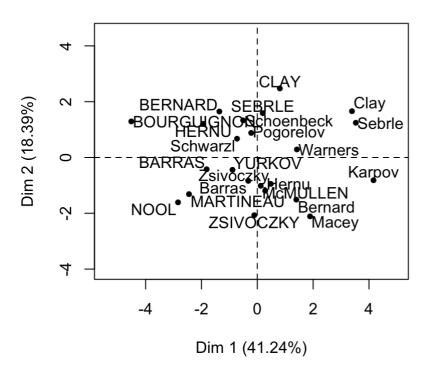
head(res.pca\$ind\$contrib)

	Dim.1	Dim.2	Dim.3	Dim.4	Dim.5
SEBRLE	0.04029447	5.9714533	1.4483919	0.03734589	8.45894063
CLAY	0.68805664	14.4839248	6.7537381	8.94458283	4.21794385
BERNARD	1.94740183	6.4234107	0.1411345	20.46819433	0.04393073
YURKOV	0.83308415	0.4632733	22.4517396	2.69663605	1.03075263
ZSIVOCZKY	0.01232413	10.1217143	6.2464325	0.05469230	0.25151025
McMULLEN	0.01549089	2.4310854	2.6102794	9.55055888	16.29493304

Graph of individuals using FactoMineR base graph

plot(res.pca, choix = "ind")

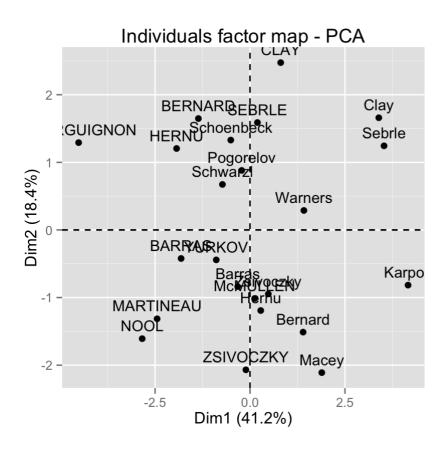
Individuals factor map (PCA)



Graph of individuals using factoextra

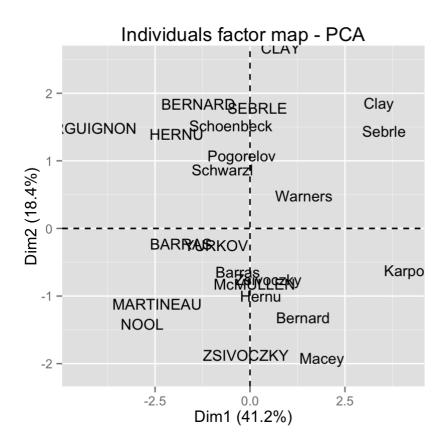
The function **fviz_pca_ind()** is used to visualize individuals :

fviz_pca_ind(res.pca)



Remove the points from the graph, use texts only:

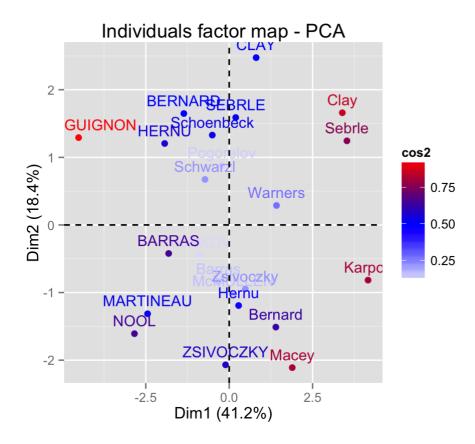
fviz_pca_ind(res.pca, geom="text")



Note that, allowed values for the argument **geom** are :

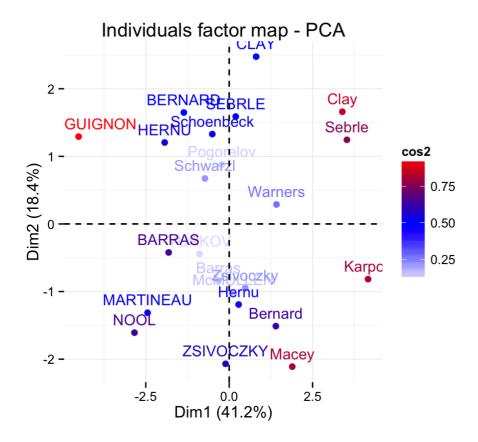
- "point" to show only points (dots)
- "text" to show only labels
- c("point", "text") to show both types

Control automatically the color of individuals using the cos2 values (the quality of the individuals on the factor map):



Read more about ggplot2 and colors here : <u>ggplot2 colors - How to change colors automatically and manually?</u>

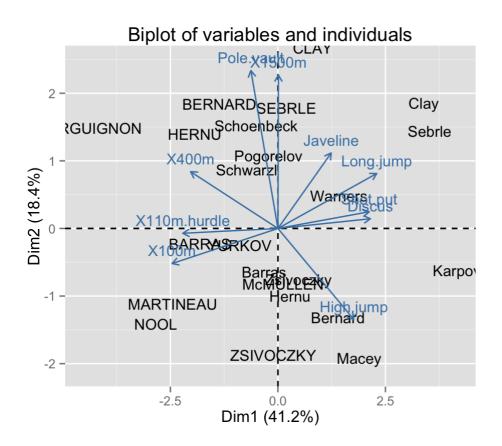
Change the theme:



Read more about ggplot2 themes here: ggplot2 themes and background colors

Make a biplot of individuals and variables:

fviz_pca_biplot(res.pca, geom = "text")



Change the color of individuals by groups

We will use iris data sets in this section:

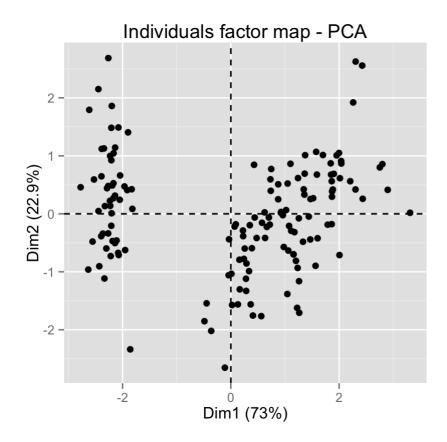
```
data(iris)
head(iris)
```

```
Sepal.Length Sepal.Width Petal.Length Petal.Width Species
1
           5.1
                        3.5
                                     1.4
                                                  0.2 setosa
2
           4.9
                       3.0
                                     1.4
                                                  0.2
                                                       setosa
3
           4.7
                        3.2
                                     1.3
                                                  0.2
                                                      setosa
4
           4.6
                        3.1
                                     1.5
                                                  0.2
                                                       setosa
5
           5.0
                       3.6
                                     1.4
                                                  0.2
                                                      setosa
           5.4
                        3.9
                                     1.7
                                                  0.4 setosa
```

```
# The variable Species (index = 5) is removed
# before PCA analysis
iris.pca <- PCA(iris[,-5], graph = FALSE)</pre>
```

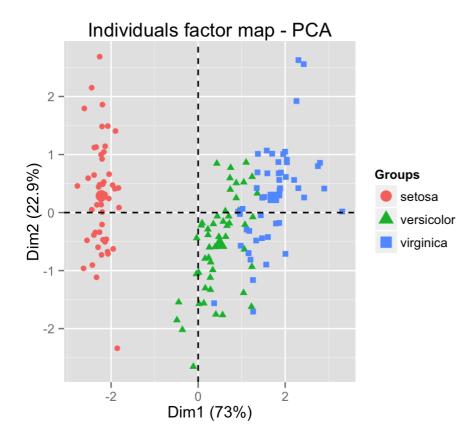
Individuals factor map:

```
# Default plot
fviz_pca_ind(iris.pca, label="none")
```

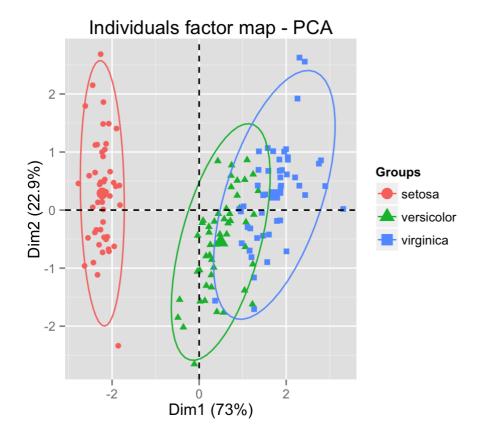


Change individual colors by groups:

fviz_pca_ind(iris.pca, label="none", habillage=iris\$Species)

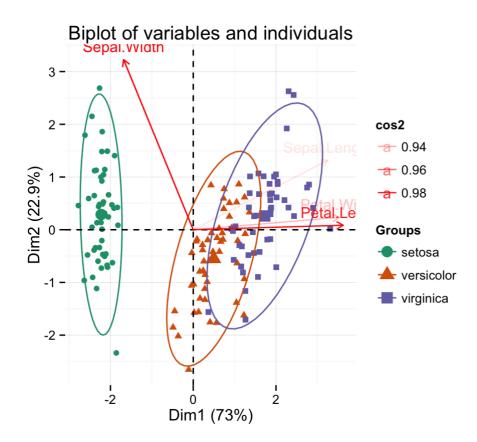


Add ellipses of point concentrations: the argument *habillage* is used to specify the factor variable for coloring the observations by groups.



- make a biplot of individuals and variables
- change the color of individuals by groups
- change the transparency of variable colors by their contribution values
- show only the labels for variables

```
fviz_pca_biplot(iris.pca,
  habillage = iris$Species, addEllipses = TRUE,
  col.var = "red", alpha.var ="cos2",
  label = "var") +
  scale_color_brewer(palette="Dark2")+
  theme_minimal()
```



Principal component analysis using supplementary individuals and variables

As described above, the data sets *decathlon2* contain **supplementary continuous variables** (quanti.sup, columns 11:12), **supplementary qualitative variables** (quali.sup, column 13) and **supplementary individuals** (ind.sup, rows 24:27)

Supplementary variables and individuals are not used for the determination of the principal components. Their coordinates are predicted using only the informations provided by the performed principal component analysis on active variables/individuals.

To specify supplementary individuals and variables, the function PCA() can be used as follow:

```
PCA(X, scale.unit = TRUE, ncp = 5, ind.sup = NULL,
    quanti.sup=NULL, quali.sup=NULL, graph=TRUE, axes = c(1,2))
```

- X: a data frame. Rows are individuals and columns are numeric variables.
- scale.unit: a logical value. If TRUE, the data are scaled to unit variance before the analysis.

- **ncp**: number of dimensions kept in the final results.
- ind.sup: a numeric vector specifying the indexes of the supplementary individuals
- quanti.sup, quali.sup: a numeric vector specifying, respectively, the indexes of the quantitative and qualitative variables
- **graph**: a logical value. If TRUE a graph is displayed.
- axes: a vector of length 2 specifying the components to be plotted

Example of usage:

Visualize supplementary quantitative variables

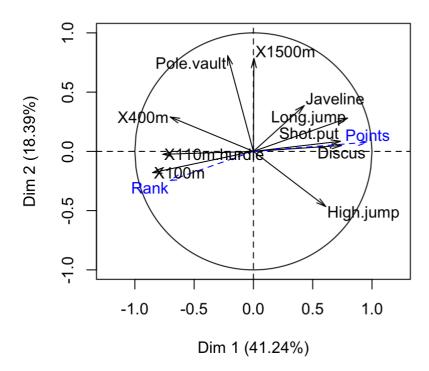
All the results (coordinates, correlation and cos2) for the supplementary quantitative variables can be extracted as follow:

```
res.pca$quanti.sup
$coord
                     Dim.2
                               Dim.3
           Dim.1
                                            Dim.4
     -0.7014777 -0.24519443 -0.1834294 0.05575186 -0.07382647
Points 0.9637075 0.07768262 0.1580225 -0.16623092 -0.03114711
$cor
           Dim.1
                      Dim.2
                                Dim.3
                                            Dim.4
Rank -0.7014777 -0.24519443 -0.1834294 0.05575186 -0.07382647
Points 0.9637075 0.07768262 0.1580225 -0.16623092 -0.03114711
$cos2
          Dim.1
                     Dim.2
                               Dim.3
                                          Dim.4
                                                      Dim.5
      0.4920710 0.060120310 0.03364635 0.00310827 0.0054503477
Points 0.9287322 0.006034589 0.02497110 0.02763272 0.0009701427
```

Variables factor map using FactoMineR base graph:

```
plot(res.pca, choix = "var")
```

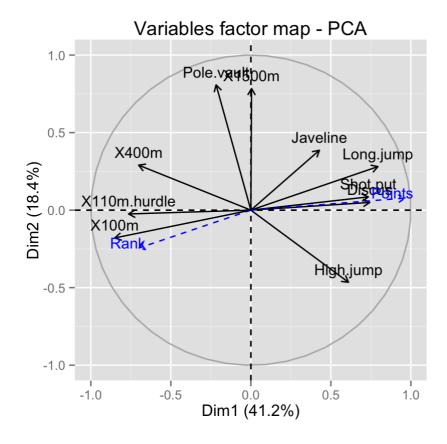
Variables factor map (PCA)



Supplementary quantitative variables are shown in blue color and dashed lines.

It's also possible to make the variables factor map using factoextra:

fviz_pca_var(res.pca)



Visualize supplementary individuals

The data sets *decathlon2* contain some **supplementary individuals** from row 24 to 27.

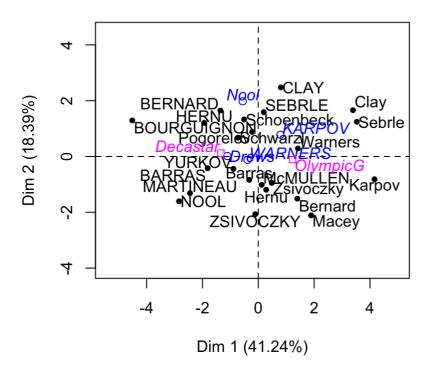
```
# Data for the supplementary individuals
ind.sup <- decathlon2[24:27, 1:10]
ind.sup[, 1:6]</pre>
```

	X100m	Long.jump	Shot.put	High.jump	X400m	X110m.hurdle
KARPOV	11.02	7.30	14.77	2.04	48.37	14.09
WARNERS	11.11	7.60	14.31	1.98	48.68	14.23
Nool	10.80	7.53	14.26	1.88	48.81	14.80
Drews	10.87	7.38	13.07	1.88	48.51	14.01

Individuals factor map using FactoMineR base graph:

plot(res.pca, choix="ind")

Individuals factor map (PCA)



Supplementary individuals are shown in blue. The levels of the supplementary qualitative variable are shown in magnenta color.

The results for supplementary individuals can be extracted as follow:

```
res.pca$ind.sup
```

```
$coord
                                           Dim.4
                                                     Dim.5
            Dim.1
                       Dim.2
                                 Dim.3
                  0.77951227 -1.6330203 1.7242283 -0.75070396
KARPOV
        0.7947206
WARNERS
        -0.3864645 -0.12159237 -1.7387332 -0.7063341 -0.03230011
                  1.97748871 -0.4830358 -2.2784526 -0.25461493
Nool
        -0.5591306
        Drews
$cos2
            Dim.1
                        Dim.2
                                  Dim.3
                                           Dim.4
KARPOV
        0.05104677\ 4.911173e-02\ 0.21553730\ 0.24028620\ 0.0455487744
WARNERS 0.02422707 2.398250e-03 0.49039677 0.08092862 0.0001692349
```

```
Nool 0.02897149 3.623868e-01 0.02162236 0.48108780 0.0060077529
Drews 0.09207094 2.269527e-05 0.69560547 0.17617609 0.0079736753

$dist
KARPOV WARNERS Nool Drews
3.517470 2.482899 3.284943 3.655527
```

Supplementary qualitative variables

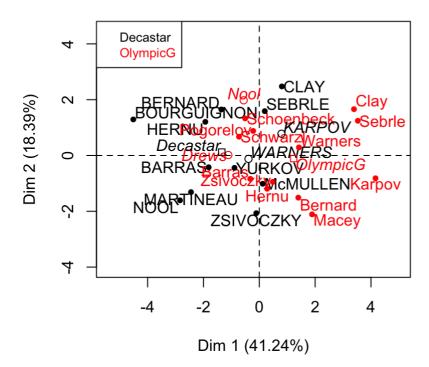
The data sets *decathlon2* contain a **supplementary qualitative variable** at columns 13 corresponding to the type of competitions.

Qualitative variable can be helpful for interpreting the data and for coloring individuals by groups.

The argument *habillage* is used to specify the index of the supplementary qualitative variable :

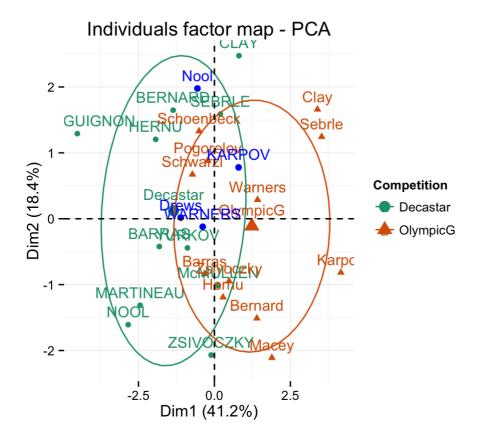
```
plot(res.pca, choix = "ind", habillage = 13)
```

Individuals factor map (PCA)



It's also possible to use factoextra:

```
fviz_pca_ind(res.pca, habillage = 13,
  addEllipses =TRUE, ellipse.level = 0.68) +
  scale_color_brewer(palette="Dark2") +
  theme_minimal()
```



Supplementary individuals are shown in blue color

The results concerning the supplementary qualitative variable are:

```
res.pca$quali
$coord
             Dim.1
                        Dim.2
                                    Dim.3
                                               Dim.4
Decastar -1.343451 0.1218097 -0.03789524
                                           0.1808357
                                                      0.1343364
OlympicG 1.231497 -0.1116589 0.03473730 -0.1657661 -0.1231417
$cos2
             Dim.1
                         Dim.2
                                      Dim.3
                                                 Dim.4
                                                              Dim.5
Decastar 0.9051233 0.007440939 0.0007201669 0.01639956 0.009050062
OlympicG 0.9051233 0.007440939 0.0007201669 0.01639956 0.009050062
$v.test
             Dim.1
                        Dim.2
                                   Dim.3
                                              Dim.4
Decastar -2.970766 0.4034256 -0.1528767
                                          0.8971036
                                                     0.7202457
OlympicG 2.970766 -0.4034256 0.1528767 -0.8971036 -0.7202457
Decastar OlympicG
1.412108 1.294433
$eta2
                           Dim 2
                                       Dim 3
                                                  Dim 4
                                                              Dim 5
Competition 0.4011568 0.00739783 0.001062332 0.03658159 0.02357972
```

Dimension description

The function **dimdesc()** can be used to identify the most correlated variables with a given principal component.

A simplified format is:

• res: an object of class PCA

• axes: a numeric vector specifying the dimensions to be described

• **prob** : the significance level

```
Example of usage:
```

```
res.desc <- dimdesc(res.pca, axes = c(1,2))
# Description of dimension 1
res.desc$Dim.1
$quanti
            correlation
                               p.value
             0.9637075 1.605675e-13
Points
             0.7941806 6.059893e-06
Long.jump
              0.7432090 4.842563e-05
Discus
          0.7339127 6.723102e-05
0.6100840 1.993677e-03
0.4282266 4.149192e-02
Shot.put
High.jump
Javeline
              -0.7014777 1.917657e-04
Rank
         -0.7016034 1.910387e-04
X400m
X110m.hurdle -0.7641252 2.195812e-05
             -0.8506257 2.727129e-07
X100m
$quali
                          p.value
                   R2
Competition 0.4011568 0.001177378
$category
          Estimate
                       p.value
OlympicG 1.287474 0.001177378
Decastar -1.287474 0.001177378
# Description of dimension 2
res.desc$Dim.2
$quanti
           correlation
                             p.value
Pole.vault 0.8074511 3.205016e-06
           0.7844802 9.384747e-06
X1500m
```

High.jump -0.4652142 2.529390e-02

Infos

This analysis has been performed using **R software** (ver. 3.1.2) and **ggplot2** (ver.)