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Principal component analysis (PCA) in R studio



JANITH PIUMAL / MARCH 18, 2023 / STATISTICS – EXPERIMENTAL DESIGN & DATA ANALYSIS USING R

```
##### Get the data#####
```

```
data("iris")
str(iris)
```

Output:

```
'data.frame':  150 obs. of  5 variables:
 $ Sepal.Length: num  5.1 4.9 4.7 4.6 5 5.4 4.6 5 4.4 4.9 ...
 $ Sepal.Width : num  3.5 3 3.2 3.1 3.6 3.9 3.4 3.4 2.9 3.1 ...
 $ Petal.Length: num  1.4 1.4 1.3 1.5 1.4 1.7 1.4 1.5 1.4 1.5 ...
 $ Petal.Width : num  0.2 0.2 0.2 0.2 0.2 0.4 0.3 0.2 0.2 0.1 ...
 $ Species      : Factor w/ 3 levels "setosa","versicolor",...: 1 1 1 1 1
1 1 1 1 1 ...
```

```
#####training datasets and test datasets building
```

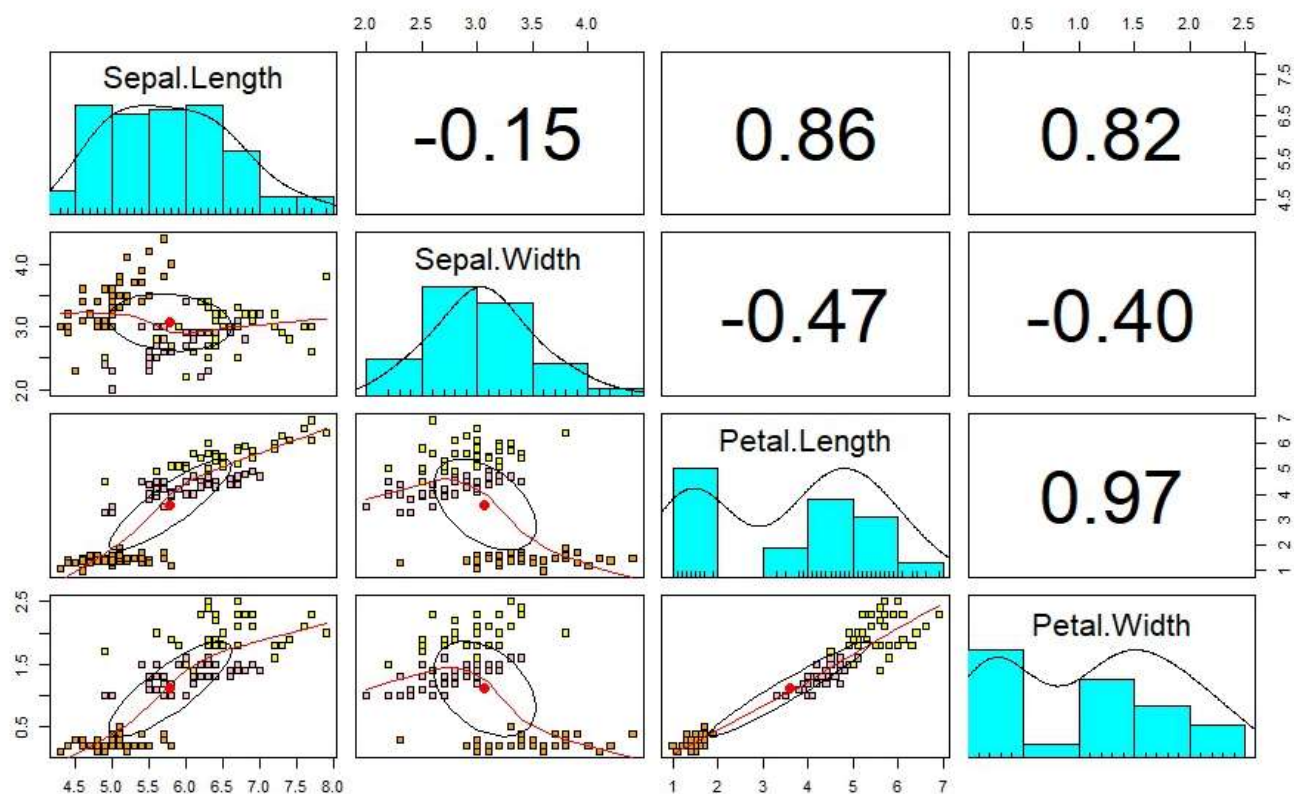
```
#####
```

```
set.seed(111)
ind <- sample(2, nrow(iris),
              replace = TRUE,
              prob = c(0.8, 0.2))
training <- iris[ind==1,]
testing <- iris[ind==2,]
```

#####Scatter Plot & Correlations#check the correlation between variables#####

```
install.packages("psych")
library(psych)
pairs.panels(training[,-5],
              gap = 1,
              bg = c("orange", "pink", "yellow")[training$Species],
              pch=22)
```

Output:



According to this petal length and petal width, sepal length and petal length , Sepal length, and petal width are highly correlated. This leads to [multicollinearity](#). This issue can be reduce using PCA analysis.

#####Principal Component Analysis#####

```
pca <- prcomp(training[,-5],
```

```

      center = TRUE,
      scale. = TRUE)
attributes(pca)

[1] "sdev"      "rotation" "center"
[4] "scale"     "x"
$class
[1] "prcomp"
pca$center
Sepal.Length Sepal.Width Petal.Length
5.8           3.1           3.6
Petal.Width
1.1

pca$scale
Sepal.Length Sepal.Width Petal.Length
0.82          0.46          1.79
Petal.Width
0.76

print(pca)

```

Output:

Standard deviations (1, .., p=4):

```
[1] 1.7173318 0.9403519 0.3843232 0.1371332
```

Rotation (n x k) = (4 x 4):

	PC1	PC2	PC3	PC4
Sepal.Length	0.5147163	-0.39817685	0.7242679	0.2279438
Sepal.Width	-0.2926048	-0.91328503	-0.2557463	-0.1220110
Petal.Length	0.5772530	-0.02932037	-0.1755427	-0.7969342
Petal.Width	0.5623421	-0.08065952	-0.6158040	0.5459403

#####summarized#####

```
summary(pca)
```

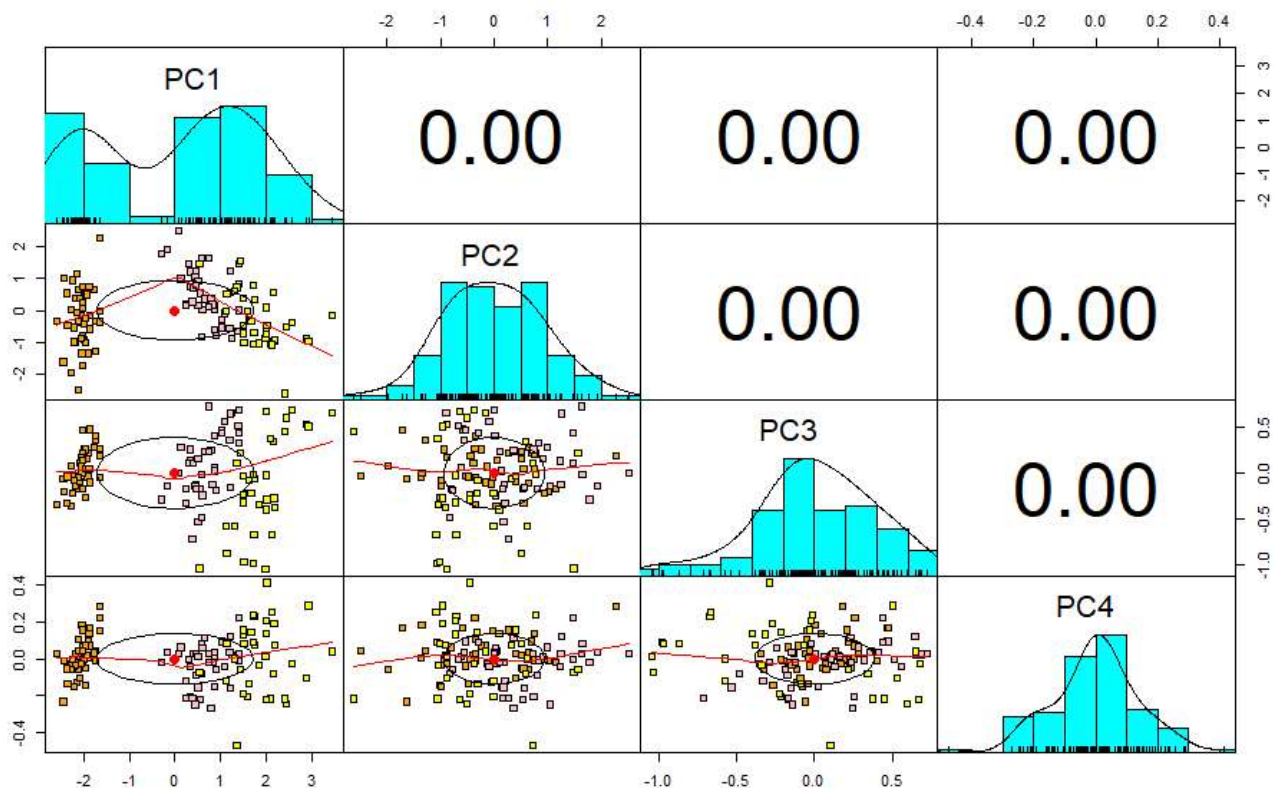
Output:

Importance of components:

	PC1	PC2	PC3	PC4
Standard deviation	1.7173	0.9404	0.38432	0.1371
Proportion of Variance	0.7373	0.2211	0.03693	0.0047
Cumulative Proportion	0.7373	0.9584	0.99530	1.0000

#####scatter plot## To check the correlation between the principal components ####

```
pairs.panels(pca$x,
              gap=0,
              bg = c("orange", "pink", "yellow")[training$Species],
              pch=22)
```

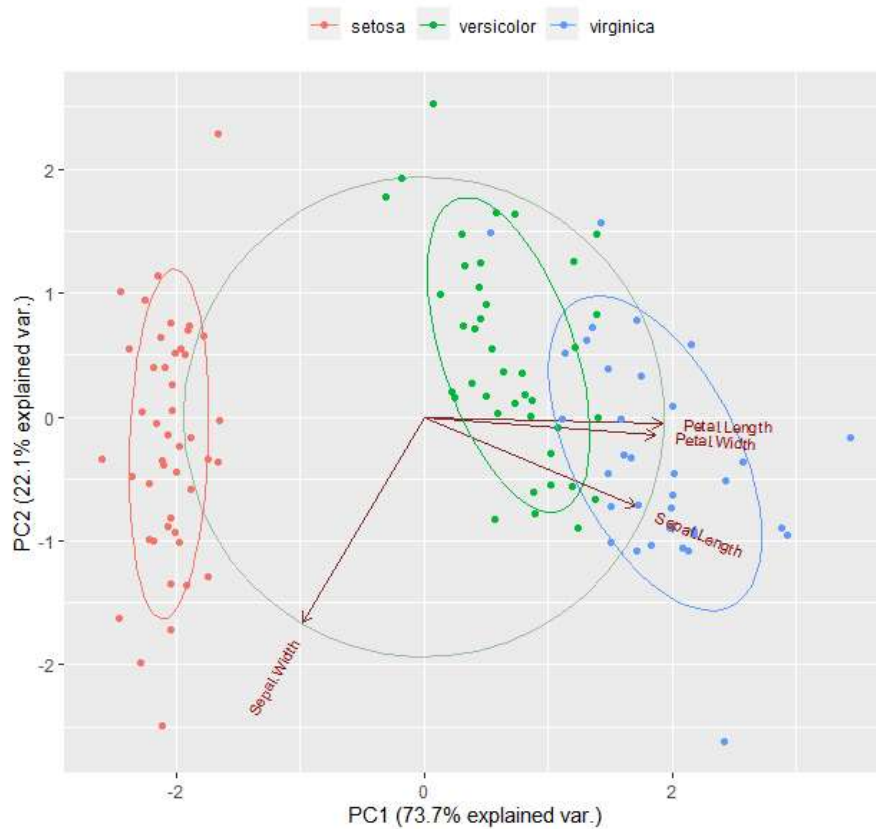
Output:

Now there is no correlation between multiple variables therefore there is no

multicollinearity issue.

```
##### explain the PCA using BY BILOT#####  
library(devtools)  
install_github("vqv/ggbiplot")  
library(ggbiplot)  
g <- ggbiplot(pca,  
              obs.scale = 1,  
              var.scale = 1,  
              groups = training$Species,  
              ellipse = TRUE,  
              circle = TRUE,  
              ellipse.prob = 0.68)  
g <- g + scale_color_discrete(name = '')  
g <- g + theme(legend.direction = 'horizontal',  
              legend.position = 'top')  
  
print(g)
```

Output:



BIPLOT is useful to understand what is happening in the data set.

- **PC1 is positively correlated with the variables Petal Length, Petal Width, Sepal Length, negatively correlated with Sepal Width.**
- **PC2 is negatively correlated with Sepal Width.**

References

- *Principal component analysis (PCA) in R | R-bloggers.* (2021, May 7). Principal Component Analysis (PCA) in R | R-bloggers. <https://www.r-bloggers.com/2021/05/principal-component-analysis-pca-in-r/>

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