

Multivariate Statistics HW2

Namwoo Kwon 20236002

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Goal : Perform the PCA of IRIS Dataset.

Process :

1. Intro 1 ~ 2
2. Quiz a - 1 : How many components do you need to adequately describe the data?
3. Quiz a - 2 : How would you interpret them?
4. Quiz b - 1 : Plot the average PC Scores for each of the three different types of iris for the first two PC
5. Quiz b - 2 : Describe your findings.

Intro 1 : Set path of file

```
setwd('/Users/namwoo/Desktop/UNIST/lecture/1-1/Multivaraiate Statistics')  
getwd()
```

```
## [1] "/Users/namwoo/Desktop/UNIST/lecture/1-1/Multivaraiate Statistics"
```

Intro 2 : Load IRIS Dataset

```
#install.packages('readxl')  
library(readxl)  
iris <- read_excel('IRIS.xlsx')
```

```
str(iris)
```

```
## tibble [150 × 5] (S3: tbl_df/tbl/data.frame)  
## $ X1: num [1:150] 1 1 1 1 1 1 1 1 1 1 ...  
## $ X2: num [1:150] 5.1 4.9 4.7 4.6 5 5.4 4.6 5 4.4 4.9 ...  
## $ X3: num [1:150] 3.5 3 3.2 3.1 3.6 3.9 3.4 3.4 2.9 3.1 ...  
## $ X4: num [1:150] 1.4 1.4 1.3 1.5 1.4 1.7 1.4 1.5 1.4 1.5 ...  
## $ X5: num [1:150] 0.2 0.2 0.2 0.2 0.2 0.4 0.3 0.2 0.2 0.1 ...
```

a - 1 : How many components do you need to adequately describe the data?

Solution Step 1 : Select Feature

```
a_iris <- iris[, 2:5]  
str(a_iris)
```

```
## tibble [150 × 4] (S3: tbl_df/tbl/data.frame)
## $ X2: num [1:150] 5.1 4.9 4.7 4.6 5 5.4 4.6 5 4.4 4.9 ...
## $ X3: num [1:150] 3.5 3 3.2 3.1 3.6 3.9 3.4 3.4 2.9 3.1 ...
## $ X4: num [1:150] 1.4 1.4 1.3 1.5 1.4 1.7 1.4 1.5 1.4 1.5 ...
## $ X5: num [1:150] 0.2 0.2 0.2 0.2 0.2 0.4 0.3 0.2 0.2 0.1 ...
```

```
summary(a_iris)
```

```
##           X2           X3           X4           X5
## Min.      :4.300   Min.      :2.000   Min.      :1.000   Min.      :0.100
## 1st Qu.:5.100   1st Qu.:2.800   1st Qu.:1.600   1st Qu.:0.300
## Median :5.800   Median :3.000   Median :4.350   Median :1.300
## Mean     :5.843   Mean     :3.057   Mean     :3.758   Mean     :1.199
## 3rd Qu.:6.400   3rd Qu.:3.300   3rd Qu.:5.100   3rd Qu.:1.800
## Max.     :7.900   Max.     :4.400   Max.     :6.900   Max.     :2.500
```

Solution Step 2 : Check Coefficient

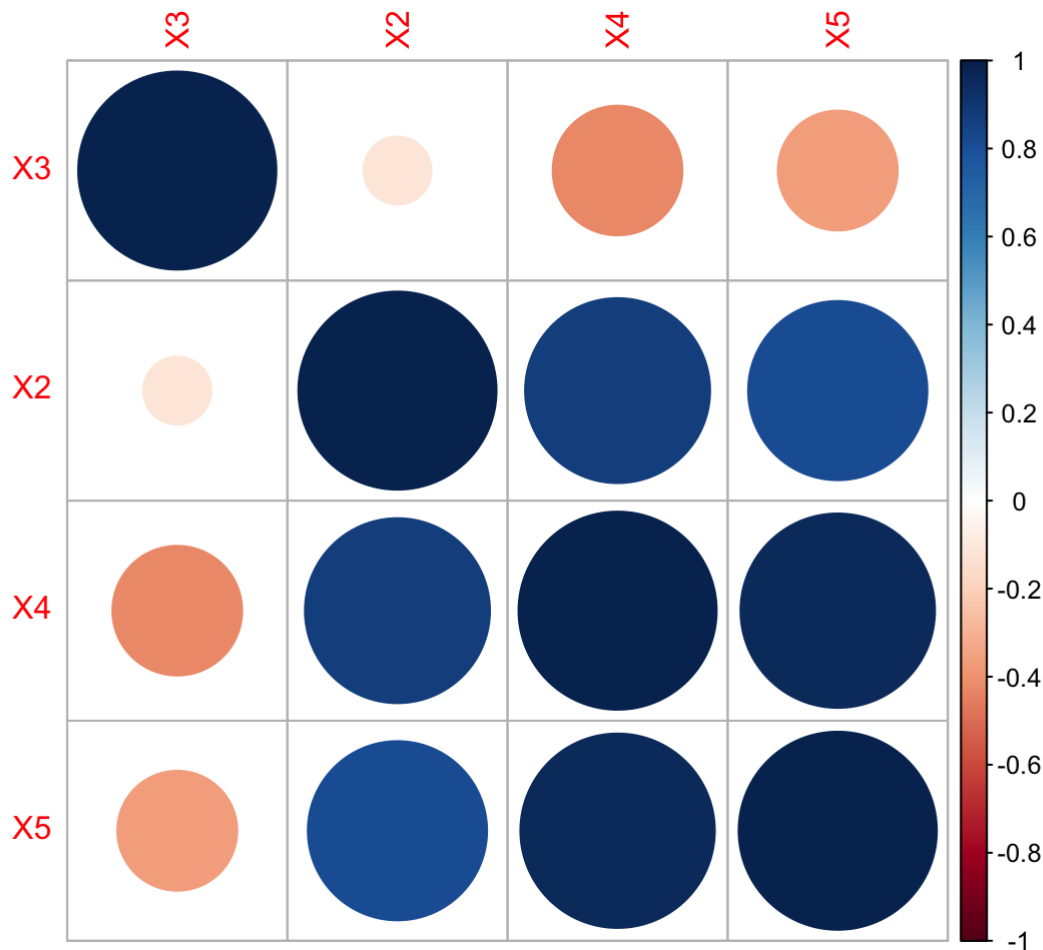
```
cor(a_iris)
```

```
##           X2           X3           X4           X5
## X2  1.0000000 -0.1175698  0.8717538  0.8179411
## X3 -0.1175698  1.0000000 -0.4284401 -0.3661259
## X4  0.8717538 -0.4284401  1.0000000  0.9628654
## X5  0.8179411 -0.3661259  0.9628654  1.0000000
```

```
#install.packages("corrplot")
library(corrplot)
```

```
## corrplot 0.92 loaded
```

```
corrplot(cor(a_iris), order="hclust")
```



Multicollinearity Issues

- You can see when variables have a high correlation between them
- EX : $\text{corr}(X2 \text{ \& } X4)$, $\text{corr}(X2 \text{ \& } X5)$
- High correlation coefficients between variables can lead to multicollinearity.

Solution Step 3 : Perform the PCA of IRIS Dataset

```
set.seed(42)

iris.pca <- prcomp(a_iris,
  # Standardization
  center=TRUE, #  $E(X_i) = 0$ 
  scale=TRUE) #  $\text{Var}(X_i) = 1$ 
```

Set PCA parameters

- 'center=TRUE' : Make the mean of all the variavles to be zero
- 'scale=TRUE' : I don't know what unit of measure each variable uses -> Need to scaled Data

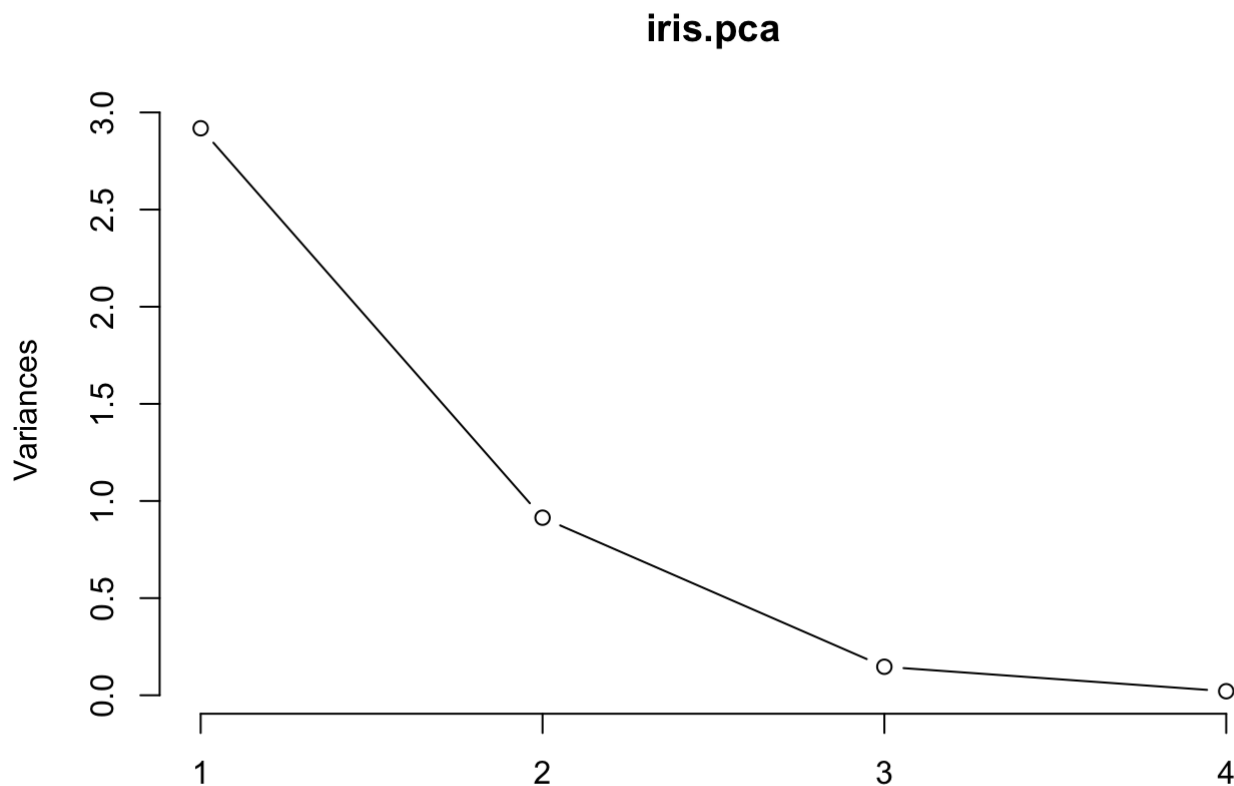
```
print(iris.pca)
```

```
## Standard deviations (1, ..., p=4):
## [1] 1.7083611 0.9560494 0.3830886 0.1439265
##
## Rotation (n x k) = (4 x 4):
##           PC1      PC2      PC3      PC4
## X2  0.5210659 -0.37741762  0.7195664  0.2612863
## X3 -0.2693474 -0.92329566 -0.2443818 -0.1235096
## X4  0.5804131 -0.02449161 -0.1421264 -0.8014492
## X5  0.5648565 -0.06694199 -0.6342727  0.5235971
```

```
summary(iris.pca)
```

```
## Importance of components:
##
##           PC1      PC2      PC3      PC4
## Standard deviation    1.7084 0.9560 0.38309 0.14393
## Proportion of Variance 0.7296 0.2285 0.03669 0.00518
## Cumulative Proportion 0.7296 0.9581 0.99482 1.00000
```

```
plot(iris.pca, type='l')
```



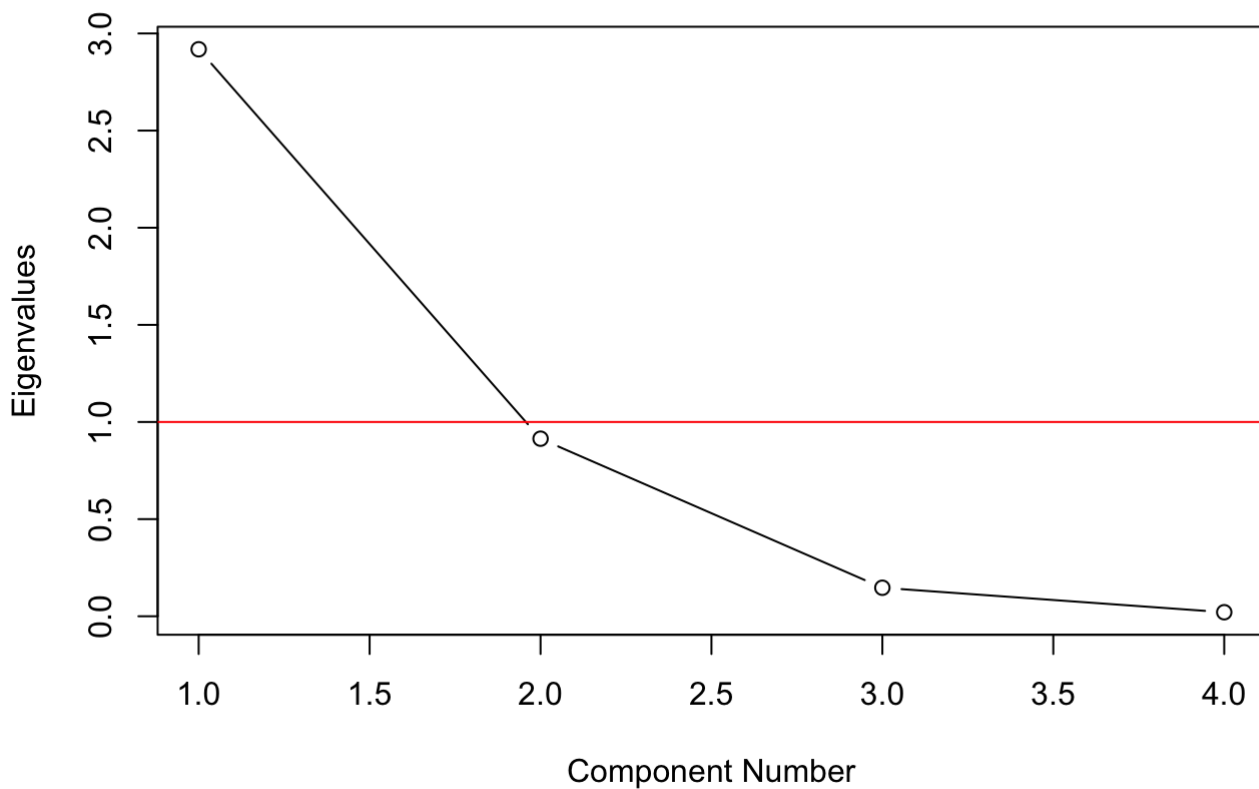
Solution Step 4 : Check Eigenvalues

```
iris.mat <- as.matrix(a_iris)
cov.mat <- cor(iris.mat)
eigen(cov.mat)
```

```
## eigen() decomposition
## $values
## [1] 2.91849782 0.91403047 0.14675688 0.02071484
##
## $vectors
##           [,1]      [,2]      [,3]      [,4]
## [1,]  0.5210659 -0.37741762  0.7195664  0.2612863
## [2,] -0.2693474 -0.92329566 -0.2443818 -0.1235096
## [3,]  0.5804131 -0.02449161 -0.1421264 -0.8014492
## [4,]  0.5648565 -0.06694199 -0.6342727  0.5235971
```

```
plot(eigen(cov.mat)$values,
     main="Scree Plot by Kaiser's Rule",
     xlab='Component Number',
     ylab='Eigenvalues',
     type='b')
abline(h=1, col='red')
```

Scree Plot by Kaiser's Rule



Answer

- It seems desirable to use 1 Principal Components by Kaiser's Rule.
- Kaiser's Rule recommends using the number of principal components with an eigenvalue greater than 1 as the final number of dimensions to use.
- Personal Suggestion : Suggest using two principal components.
 - Using only one principal component seems insufficient to explain the variance of the original data. (72.96 %)

- However, using two principal components can explain a significant portion of the variance in the original data (95.81%)

a - 2 : How would you interpret them?

Solution Step 5 : Check Principal Component Scores

```
iris.pca$x[,1:2]
```

##		PC1	PC2
##	[1,]	-2.25714118	-0.478423832
##	[2,]	-2.07401302	0.671882687
##	[3,]	-2.35633511	0.340766425
##	[4,]	-2.29170679	0.595399863
##	[5,]	-2.38186270	-0.644675659
##	[6,]	-2.06870061	-1.484205297
##	[7,]	-2.43586845	-0.047485118
##	[8,]	-2.22539189	-0.222403002
##	[9,]	-2.32684533	1.111603700
##	[10,]	-2.17703491	0.467447569
##	[11,]	-2.15907699	-1.040205867
##	[12,]	-2.31836413	-0.132633999
##	[13,]	-2.21104370	0.726243183
##	[14,]	-2.62430902	0.958296347
##	[15,]	-2.19139921	-1.853846555
##	[16,]	-2.25466121	-2.677315230
##	[17,]	-2.20021676	-1.478655729
##	[18,]	-2.18303613	-0.487206131
##	[19,]	-1.89223284	-1.400327567
##	[20,]	-2.33554476	-1.124083597
##	[21,]	-1.90793125	-0.407490576
##	[22,]	-2.19964383	-0.921035871
##	[23,]	-2.76508142	-0.456813301
##	[24,]	-1.81259716	-0.085272854
##	[25,]	-2.21972701	-0.136796175
##	[26,]	-1.94532930	0.623529705
##	[27,]	-2.04430277	-0.241354991
##	[28,]	-2.16133650	-0.525389422
##	[29,]	-2.13241965	-0.312172005
##	[30,]	-2.25769799	0.336604248
##	[31,]	-2.13297647	0.502856075
##	[32,]	-1.82547925	-0.422280389
##	[33,]	-2.60621687	-1.787587272
##	[34,]	-2.43800983	-2.143546796
##	[35,]	-2.10292986	0.458665270
##	[36,]	-2.20043723	0.205419224
##	[37,]	-2.03831765	-0.659349230
##	[38,]	-2.51889339	-0.590315163
##	[39,]	-2.42152026	0.901161067
##	[40,]	-2.16246625	-0.267981199
##	[41,]	-2.27884081	-0.440240541
##	[42,]	-1.85191836	2.329610745
##	[43,]	-2.54511203	0.477501017
##	[44,]	-1.95788857	-0.470749613
##	[45,]	-2.12992356	-1.138415464
##	[46,]	-2.06283361	0.708678586
##	[47,]	-2.37677076	-1.116688691
##	[48,]	-2.38638171	0.384957230
##	[49,]	-2.22200263	-0.994627669
##	[50,]	-2.19647504	-0.009185585
##	[51,]	1.09810244	-0.860091033
##	[52,]	0.72889556	-0.592629362
##	[53,]	1.23683580	-0.614239894
##	[54,]	0.40612251	1.748546197

```
## [55,] 1.07188379 0.207725147
## [56,] 0.38738955 0.591302717
## [57,] 0.74403715 -0.770438272
## [58,] -0.48569562 1.846243998
## [59,] 0.92480346 -0.032118478
## [60,] 0.01138804 1.030565784
## [61,] -0.10982834 2.645211115
## [62,] 0.43922201 0.063083852
## [63,] 0.56023148 1.758832129
## [64,] 0.71715934 0.185602819
## [65,] -0.03324333 0.437537419
## [66,] 0.87248429 -0.507364239
## [67,] 0.34908221 0.195656268
## [68,] 0.15827980 0.789451008
## [69,] 1.22100316 1.616827281
## [70,] 0.16436725 1.298259939
## [71,] 0.73521959 -0.395247446
## [72,] 0.47469691 0.415926887
## [73,] 1.23005729 0.930209441
## [74,] 0.63074514 0.414997441
## [75,] 0.70031506 0.063200094
## [76,] 0.87135454 -0.249956017
## [77,] 1.25231375 0.076998069
## [78,] 1.35386953 -0.330205463
## [79,] 0.66258066 0.225173502
## [80,] -0.04012419 1.055183583
## [81,] 0.13035846 1.557055553
## [82,] 0.02337438 1.567225244
## [83,] 0.24073180 0.774661195
## [84,] 1.05755171 0.631726901
## [85,] 0.22323093 0.286812663
## [86,] 0.42770626 -0.842758920
## [87,] 1.04522645 -0.520308714
## [88,] 1.04104379 1.378371048
## [89,] 0.06935597 0.218770433
## [90,] 0.28253073 1.324886147
## [91,] 0.27814596 1.116288852
## [92,] 0.62248441 -0.024839814
## [93,] 0.33540673 0.985103828
## [94,] -0.36097409 2.012495825
## [95,] 0.28762268 0.852873116
## [96,] 0.09105561 0.180587142
## [97,] 0.22695654 0.383634868
## [98,] 0.57446378 0.154356489
## [99,] -0.44617230 1.538637456
## [100,] 0.25587339 0.596852285
## [101,] 1.83841002 -0.867515056
## [102,] 1.15401555 0.696536401
## [103,] 2.19790361 -0.560133976
## [104,] 1.43534213 0.046830701
## [105,] 1.86157577 -0.294059697
## [106,] 2.74268509 -0.797736709
## [107,] 0.36579225 1.556289178
## [108,] 2.29475181 -0.418663020
## [109,] 1.99998633 0.709063226
## [110,] 2.25223216 -1.914596301
```



```
## [111,] 1.35962064 -0.690443405
## [112,] 1.59732747 0.420292431
## [113,] 1.87761053 -0.417849815
## [114,] 1.25590769 1.158379741
## [115,] 1.46274487 0.440794883
## [116,] 1.58476820 -0.673986887
## [117,] 1.46651849 -0.254768327
## [118,] 2.41822770 -2.548124795
## [119,] 3.29964148 -0.017721580
## [120,] 1.25954707 1.701046715
## [121,] 2.03091256 -0.907427443
## [122,] 0.97471535 0.569855257
## [123,] 2.88797650 -0.412259950
## [124,] 1.32878064 0.480202496
## [125,] 1.69505530 -1.010536476
## [126,] 1.94780139 -1.004412720
## [127,] 1.17118007 0.315338060
## [128,] 1.01754169 -0.064131184
## [129,] 1.78237879 0.186735633
## [130,] 1.85742501 -0.560413289
## [131,] 2.42782030 -0.258418706
## [132,] 2.29723178 -2.617554417
## [133,] 1.85648383 0.177953334
## [134,] 1.11042770 0.291944582
## [135,] 1.19845835 0.808606364
## [136,] 2.78942561 -0.853942542
## [137,] 1.57099294 -1.065013214
## [138,] 1.34179696 -0.421020154
## [139,] 0.92173701 -0.017165594
## [140,] 1.84586124 -0.673870645
## [141,] 2.00808316 -0.611835930
## [142,] 1.89543421 -0.687273065
## [143,] 1.15401555 0.696536401
## [144,] 2.03374499 -0.864624030
## [145,] 1.99147547 -1.045665670
## [146,] 1.86425786 -0.385674038
## [147,] 1.55935649 0.893692855
## [148,] 1.51609145 -0.268170747
## [149,] 1.36820418 -1.007877934
## [150,] 0.95744849 0.024250427
```

```
pcscores<-data.frame(iris.pca$x[,1:2])
pcscores$species<-iris$X1
head(pcscores)
```

```
##      PC1      PC2 species
## 1 -2.257141 -0.4784238      1
## 2 -2.074013  0.6718827      1
## 3 -2.356335  0.3407664      1
## 4 -2.291707  0.5953999      1
## 5 -2.381863 -0.6446757      1
## 6 -2.068701 -1.4842053      1
```

```
# Calculate the average PC score for each Iris species
iris.pca.mean <- aggregate (~species , data=pcscores , mean)
iris.pca.mean
```

```
##      species      PC1      PC2
## 1         1 -2.2173249 -0.2879627
## 2         2  0.4947904  0.5483335
## 3         3  1.7225345 -0.2603708
```

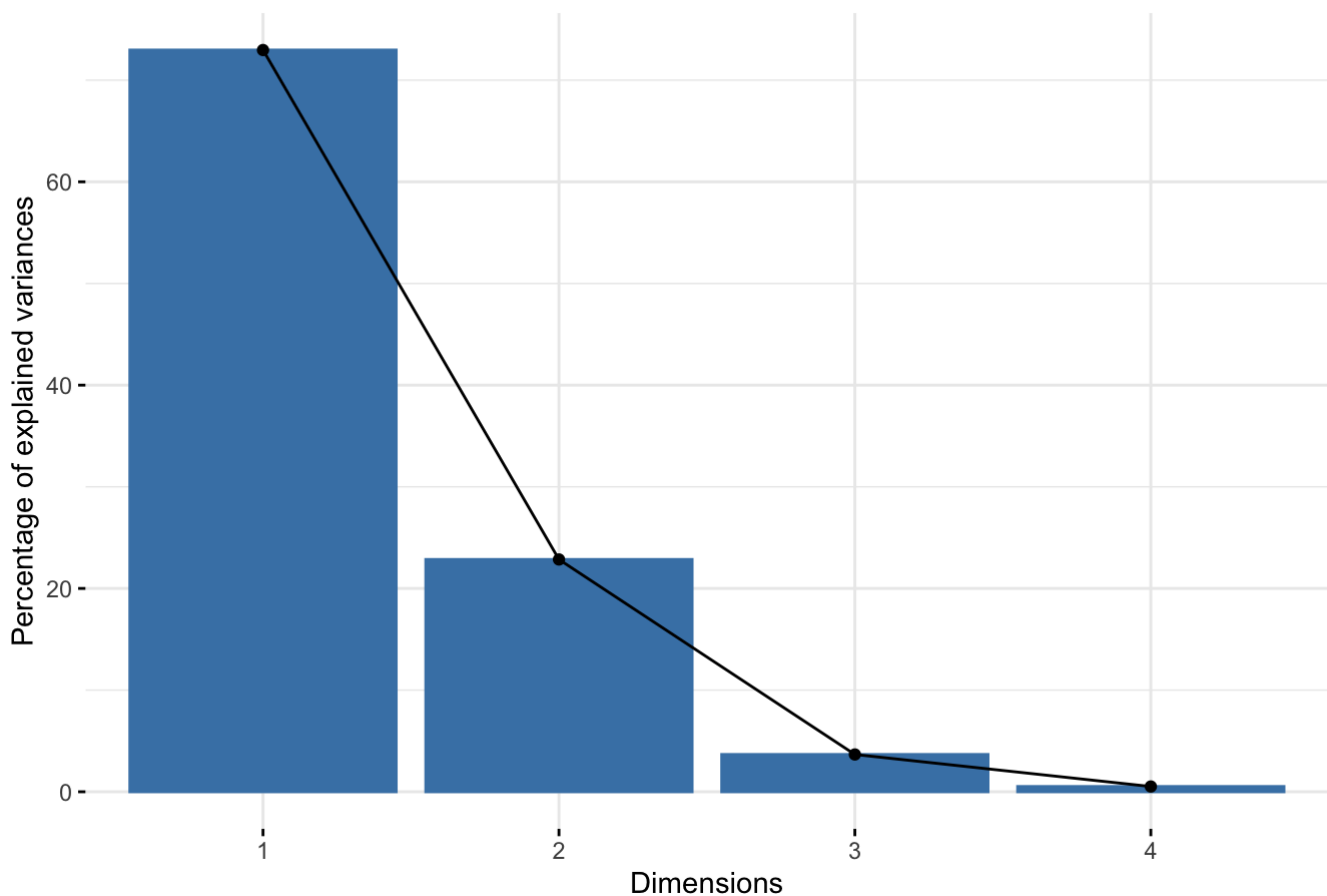
```
#install.packages('factoextra')
library(factoextra)
```

```
## Loading required package: ggplot2
```

```
## Welcome! Want to learn more? See two factoextra-related books at https://goo.gl/ve3WBa
```

```
fviz_eig(iris.pca)
```

Scree plot

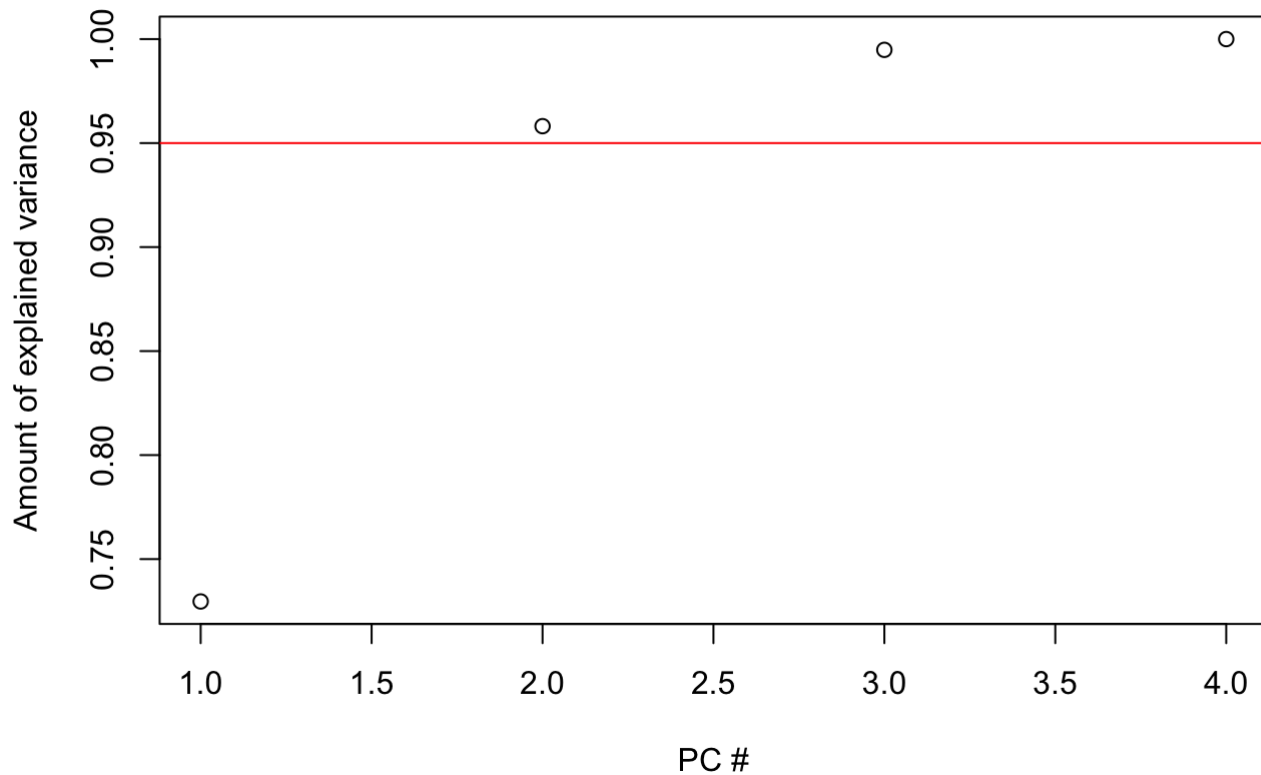


Percentage of explained variance by Number of Dimensions

- The first through second dimensions explain at least 25% of the variance in the original data.

```
p<-length(iris.pca$sdev)
cumpro <- cumsum(iris.pca$sdev^2 / sum(iris.pca$sdev^2))
plot(cumpro[0:p], xlab = "PC #", ylab = "Amount of explained variance", main = "Cumulative variance plot")
abline(h=0.95, col='red')
```

Cumulative variance plot

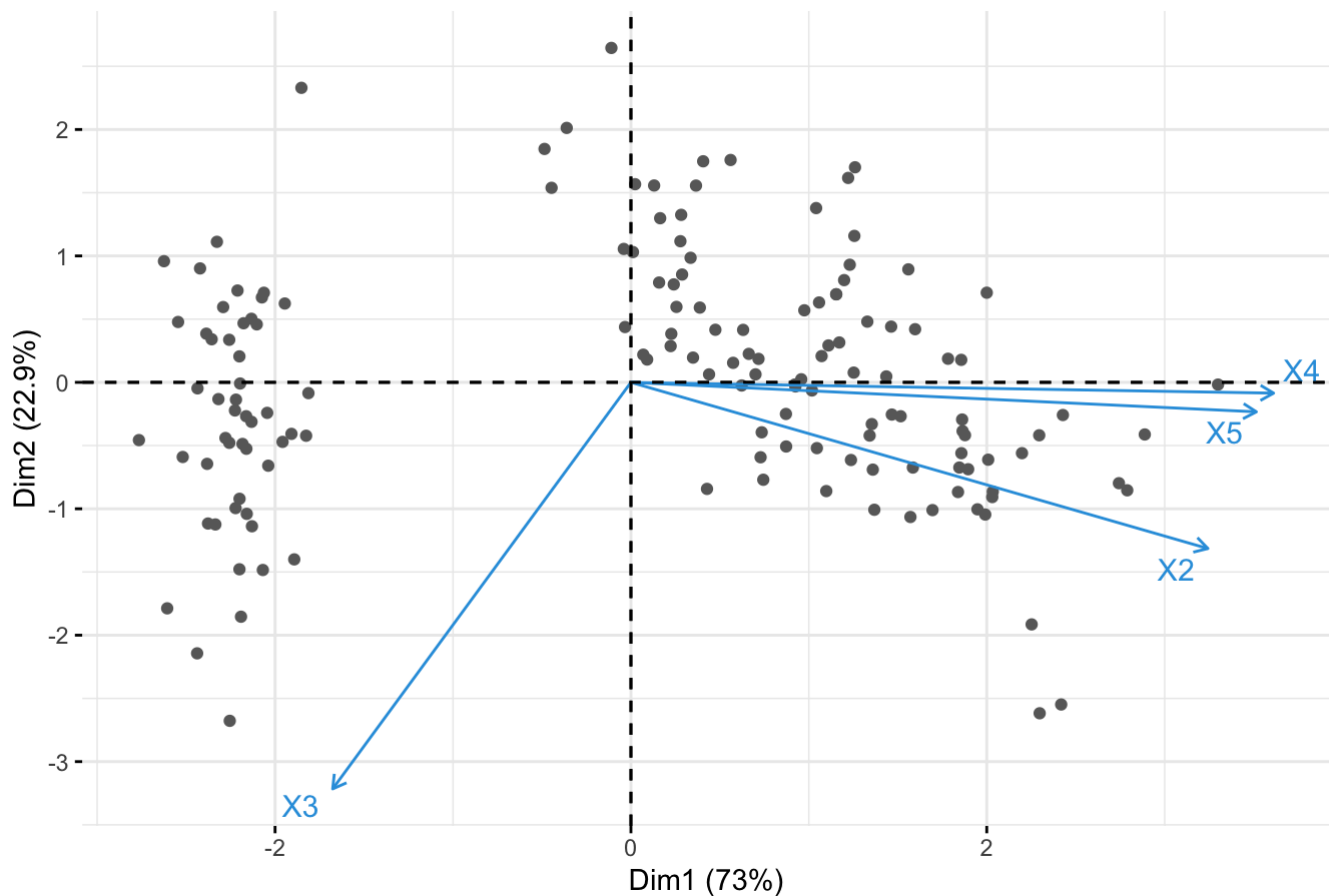


Cumulative Amount of explained variance by Number of Dimensions

- Just two dimensions can explain more than 95% of the variance in your source data.

```
fviz_pca_biplot(iris.pca, geom = "point", repel = TRUE,
                 col.var = "#2E9FDF", # Variables color
                 col.ind = "#696969"  # Individuals color
)
```

PCA - Biplot



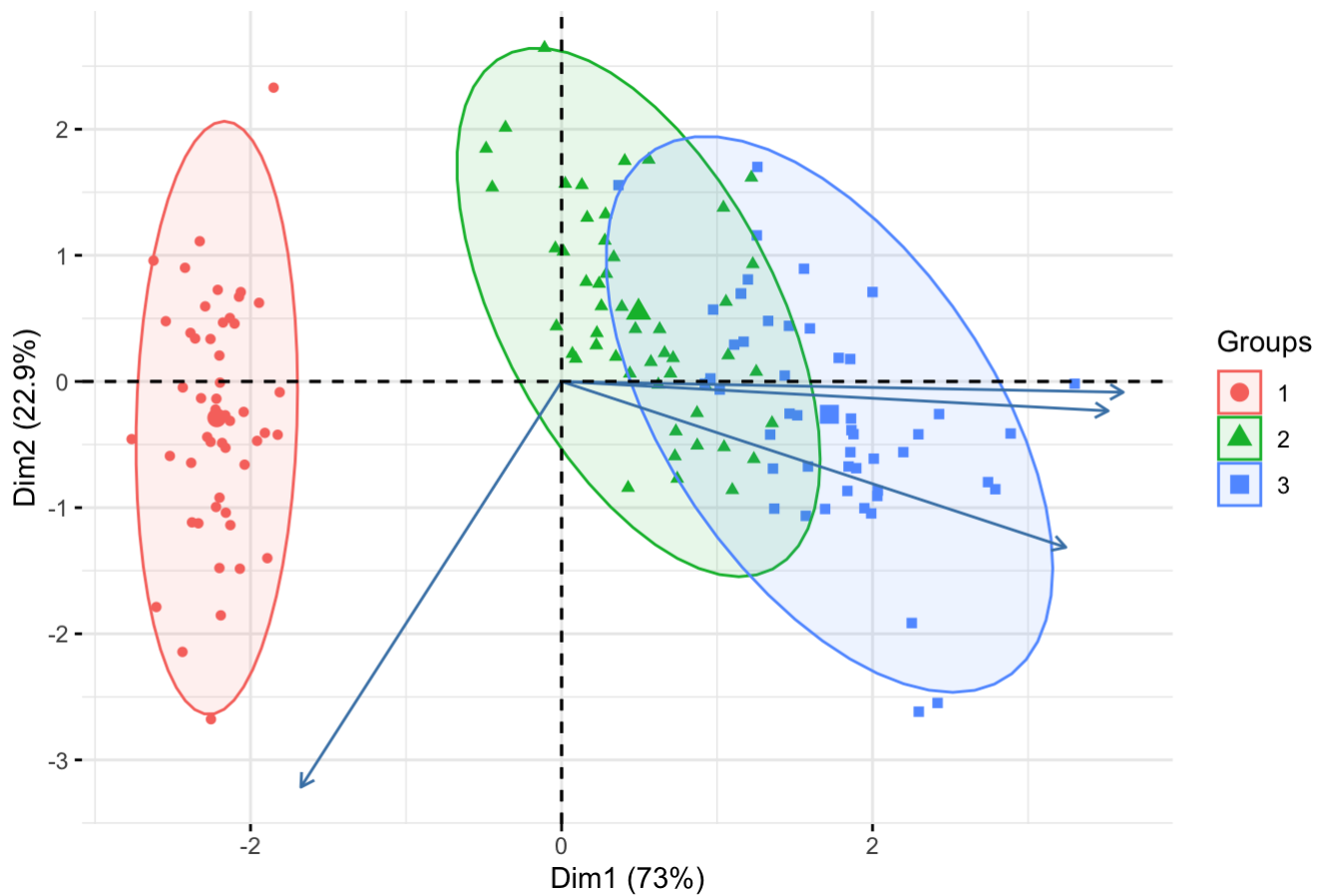
Explain the relationship between PC and source data variables with Biplot

- The variables that had the most impact on PC1 are X4, X5 and X2 (In order of greatest impact)
- The variables that had the most impact on PC2 are X3

b - 1 : Plot the average PC Scores for each of the three different types of iris for the first two PC

```
fviz_pca_biplot(iris.pca,
                 label='none',
                 habillage=iris$X1,
                 addEllipses = TRUE,
                 palette = 'jco',
                 )
```

PCA - Biplot



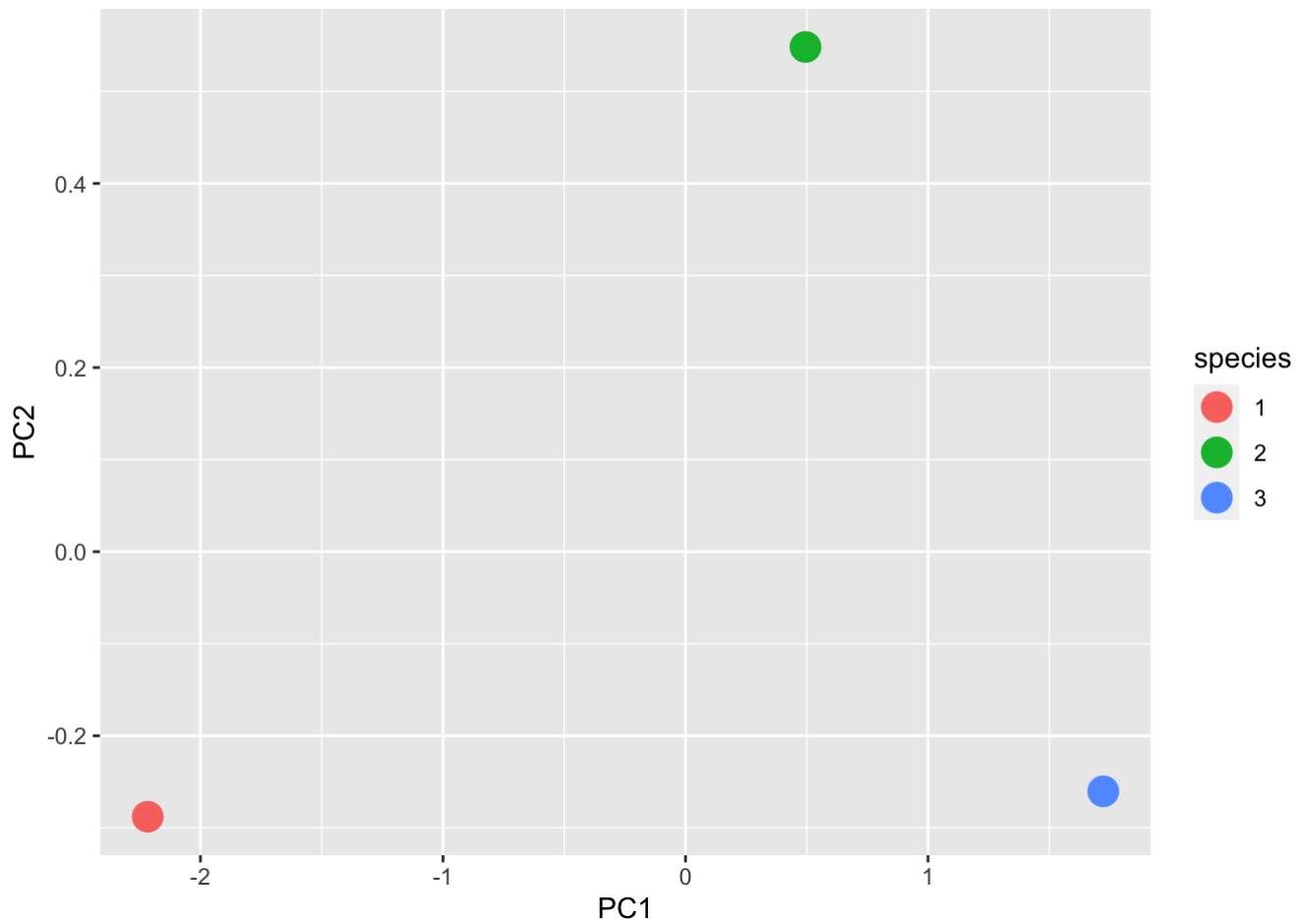
```
library("ggplot2")
```

```
iris.pca.mean
```

```
## species      PC1      PC2
## 1      1 -2.2173249 -0.2879627
## 2      2  0.4947904  0.5483335
## 3      3  1.7225345 -0.2603708
```

```
iris.pca.mean$species = as.character(iris.pca.mean$species)
```

```
ggplot(data=iris.pca.mean, aes(x=PC1, y=PC2, color=species)) + geom_point(size=5)
```



b - 2 : Describe your findings.

- Based on PC1, it becomes easier to distinguish Specie 1 from the rest.
- Based on PC2, it becomes easier to distinguish Specie 2 from the rest
- Drawing a diagonal line between PC1 and PC2 makes it easier to distinguish PC3 from the rest.