# TMA4268 Statistical Learning

Chapter 10: Unsupervised Learning - Lab 1

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# Lab 1: Principal Components Analysis

We are going to use the USArrests dataset. It contains crime statistics per 100000 residents in 50 states of the USA. The crime types included are Assault, Murder and Rape. In addition, it contains information about the percent of the population in each state living in urban areas.

## **Data Exploration**

#### Data rows

The rows of the data contains the 50 states, in alphabetical order.

```
states=row.names(USArrests) # part of the base R package states
```

```
[1] "Alabama"
                          "Alaska"
                                             "Arizona"
                                                               "Arkansas"
##
    [5] "California"
                          "Colorado"
                                             "Connecticut"
                                                               "Delaware"
   [9] "Florida"
                          "Georgia"
                                             "Hawaii"
                                                               "Idaho"
## [13] "Illinois"
                          "Indiana"
                                             "Iowa"
                                                               "Kansas"
## [17] "Kentucky"
                          "Louisiana"
                                             "Maine"
                                                               "Maryland"
## [21] "Massachusetts"
                          "Michigan"
                                             "Minnesota"
                                                               "Mississippi"
       "Missouri"
                          "Montana"
                                             "Nebraska"
                                                               "Nevada"
## [29] "New Hampshire"
                          "New Jersey"
                                             "New Mexico"
                                                               "New York"
## [33]
        "North Carolina"
                          "North Dakota"
                                             "Ohio"
                                                               "Oklahoma"
## [37]
        "Oregon"
                                             "Rhode Island"
                                                               "South Carolina"
                          "Pennsylvania"
                                                               "Utah"
## [41] "South Dakota"
                          "Tennessee"
                                             "Texas"
## [45] "Vermont"
                          "Virginia"
                                             "Washington"
                                                               "West Virginia"
                          "Wyoming"
   [49] "Wisconsin"
```

#### Data columns

The columns of the data contain the four variables

```
names(USArrests)
```

```
## [1] "Murder" "Assault" "UrbanPop" "Rape"
```

#### Mean and variance

The variables have vastly different means

```
apply(USArrests, 2, mean)

## Murder Assault UrbanPop Rape
## 7.788 170.760 65.540 21.232

The same is true for the variances
apply(USArrests, 2, var)
```

```
## Murder Assault UrbanPop Rape
## 18.97047 6945.16571 209.51878 87.72916
```

If we failed to scale the variables before performing PCA, then most of the principal components that we observed would be driven by the Assault variable, since it has by far the largest mean and variance.

#### **PCA**

We now go on to apply PCA on the standardized variables (mean 0 and std. dev. 1). By default, the prcomp() function centers the variables to have mean zero. By using the option scale=TRUE, we scale the variables to have standard deviation one.

```
pr.out=prcomp(USArrests, scale=TRUE)
```

## PCA output

The output from prcomp() contains a number of useful quantities.

```
names(pr.out)
## [1] "sdev" "rotation" "center" "scale" "x"
```

center and scale The center and scale components correspond to the means and standard deviations of the variables that were used for scaling prior to implementing PCA.

```
pr.out$center
```

```
## Murder Assault UrbanPop Rape
## 7.788 170.760 65.540 21.232
pr.out$scale
```

```
## Murder Assault UrbanPop Rape
## 4.355510 83.337661 14.474763 9.366385
```

rotation The rotation matrix provides the principal component loadings; each column of pr.out\$rotation contains the corresponding principal component loading vector.

```
pr.out$rotation
```

```
## PC1 PC2 PC3 PC4

## Murder -0.5358995 0.4181809 -0.3412327 0.64922780

## Assault -0.5831836 0.1879856 -0.2681484 -0.74340748

## UrbanPop -0.2781909 -0.8728062 -0.3780158 0.13387773

## Rape -0.5434321 -0.1673186 0.8177779 0.08902432
```

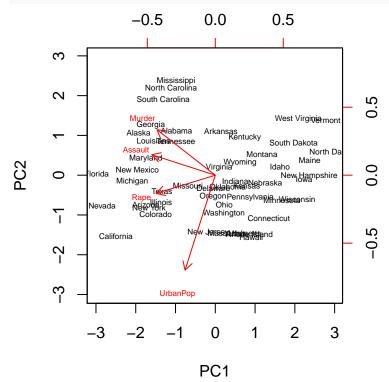
**Score vectors** The  $50 \times 4$  matrix **x** has as its columns the principal component score vectors. That is, the kth column is the k-th principal component score vector.

```
dim(pr.out$x)
```

## [1] 50 4

**Biplot** We can plot the first two principal components as follows:

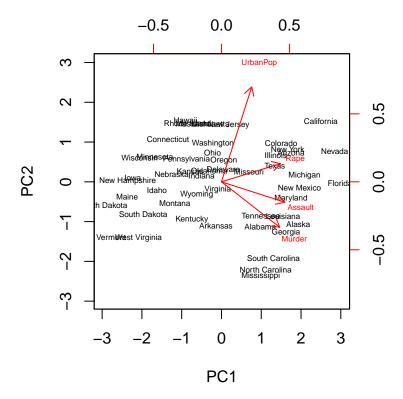
```
biplot(pr.out, scale=0, cex = 0.5)
```



The scale=0 argument to biplot() ensures that the arrows are scaled to represent the loadings; other values for scale give slightly different biplots with different interpretations.

Remember that the PCs are unique up to a sign change, so the code below should give equivalent results

```
pr.out$rotation=-pr.out$rotation
pr.out$x=-pr.out$x
biplot(pr.out, scale=0, cex = 0.5)
```



**Proportion of variance explained (PVE)** The prcomp() function also outputs the standard deviation of each principal component.

```
pr.out$sdev
```

### ## [1] 1.5748783 0.9948694 0.5971291 0.4164494

The variance explained by each principal component is obtained by squaring these:

```
pr.var=pr.out$sdev^2
pr.var
```

## ## [1] 2.4802416 0.9897652 0.3565632 0.1734301

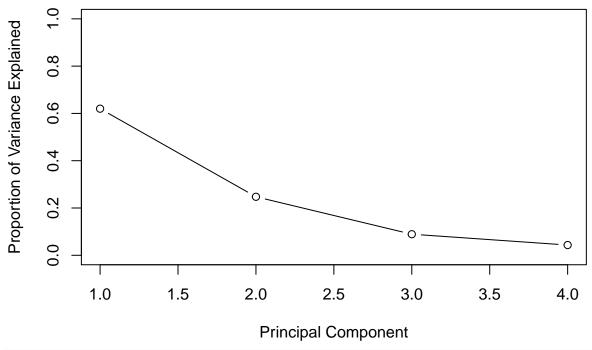
We can then compute the proportion of variance explained by each principal component

```
pve=pr.var/sum(pr.var)
pve
```

## ## [1] 0.62006039 0.24744129 0.08914080 0.04335752

We can plot the PVE explained by each component, as well as the cumulative PVE, as follows:

```
plot(pve,
     xlab="Principal Component",
     ylab="Proportion of Variance Explained",
     ylim=c(0,1),
     type='b')
```



```
plot(cumsum(pve),
    xlab="Principal Component",
    ylab="Cumulative Proportion of Variance Explained",
    ylim=c(0,1),
    type='b')
```

