Chapter 7: Principal Component Analysis

Assumes that  $X=(x_{ij})$  is normalised with  $\frac{1}{n}\sum_{i=1}^n x_{ij}=0$  and  $\frac{1}{n}\sum_{i=1}^n x_{ij}^2=1$ . We look for the 1st PC that ensures linear combination of the sample feature values  $z_{i1}=\phi_{11}x_{i1}+\cdots+\phi_{p1}x_{ip}$  has largest sample variance. In other words, we seek for  $\phi_{i1}, j=1,\ldots,p$ :

$$\min_{\phi_{11},...,\phi_{p1}} \left\{ \frac{1}{n} \sum_{i=1}^{n} \left( \sum_{j=1}^{p} \phi_{j1} x_{ij} \right)^{2} \right\} \quad s.t. \sum_{j=1}^{p} \phi_{j1}^{2} = 1,$$

Then, step by step, look for the 2nd, 3rd, ..., p-th PC that also has the largest sample variance and linear independent with all the previous PC's:

$$\min_{\phi_{12},\ldots,\phi_{p2}} \left\{ \frac{1}{n} \sum_{i=1}^{n} \left( \sum_{j=1}^{p} \phi_{j2} x_{ij} \right)^{2} \right\} \quad s.t. \sum_{j=1}^{p} \phi_{j2}^{2} = 1, \sum_{j=1}^{p} \phi_{j2} \phi_{j1} = 0,$$

$$\min_{\phi_{13},...,\phi_{p3}} \left\{ \frac{1}{n} \sum_{i=1}^{n} \left( \sum_{j=1}^{p} \phi_{j3} x_{ij} \right)^{2} \right\} \quad s.t. \sum_{j=1}^{p} \phi_{j3}^{2} = 1, \sum_{j=1}^{p} \phi_{j3} \phi_{j1} = 0, \sum_{j=1}^{p} \phi_{j3} \phi_{j2} = 0,$$

et al.

We perform PCA on the "USArrests" data set. The row of the data set contain 50 states, in alphabetical order.

```
L > states <-row.names(USArrests)
> states
```

The column of the data contain the four variables.

```
> names(USArrests)
2  [1] "Murder" "Assault" "UrbanPop" "Rape"
> apply(USArrests, 2, var)
4  Murder Assault UrbanPop Rape
18.97047 6945.16571 209.51878 87.72916
```

4 variables have vastly different means and variance, hence it is important to standardise the variables to have mean zero and std one before performing PCA.

prcomp() function is used to perform PCA. By default, the prcomp() function centres the variables to have mean zero. By using the option scale=TRUE, we scale the variables to have deviation one.

```
1 > pr_out <-prcomp(USArrests, scale = TRUE)
> names(pr_out)
3 [1] "sdev" "rotation" "center" "scale" "x"
```

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

```
1 > 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
```

The rotation matrix provides the principal component loadings, each column contains the corresponding PCA loading vector.

```
> pr_out$rotation

PC1 PC2 PC3 PC4

Murder -0.5358995 0.4181809 -0.3412327 0.64922780

4 Assault -0.5831836 0.1879856 -0.2681484 -0.74340748

UrbanPop -0.2781909 -0.8728062 -0.3780158 0.13387773

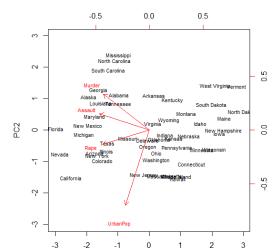
6 Rape -0.5434321 -0.1673186 0.8177779 0.08902432
```

The 50 by 4 matrix  $\times$  has its columns the principal component score vectors, i.e. the k-th column is the k-th PC score vector.

```
> dim(pr_out$x)
  [1] 50 4
 > head(pr_out$x, 5)
                    PC1
                               PC2
                                           PC3
                                                      PC4
4
  Alabama
             -0.9756604
                       1 1220012 -0 43980366
                                                0 1546966
 Alaska
            -1.9305379 1.0624269
                                    2.01950027 - 0.4341755
  Arizona
          -1.7454429 -0.7384595
                                    0.05423025 - 0.8262642
 Arkansas 0.1399989
                       1.1085423
                                    0.11342217 - 0.1809736
  California -2.4986128 -1.5274267
                                    0.59254100 - 0.3385592
```

We can plot the first two principal components as follows. The scale=0 argument to biplot() ensures that the arrows are scaled to represent the loadings.

```
1 > biplot(pr_out, scale=0, cex=0.7)
```



The summary() function for prcomp() outputs show the standard deviation, proportion of variance explained (PVE) of each principal component, and the cumulative PVE.

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

```
par(mfrow = c(1, 2))
  pr_imp <- summary(pr_out)$importance</pre>
  plot(pr_imp[2,], xlab="Principal Component",
       ylab="PVE", ylim=c(0,1), type='b')
  plot(pr_imp[3,], xlab="Principal Component",
       ylab="Cumulative PVE", ylim=c(0,1), type='b')
6
  par(mfrow = c(1, 1))
```

