14) Principal Components Analysis

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March 2019

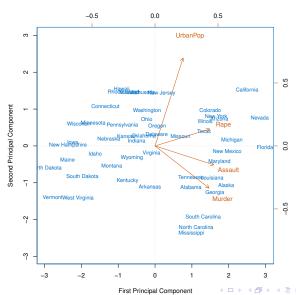
Reference

Tables, Graphics, and Figures from

James et al. (2017): Ch 10.2, and 10.4

Hastie et al. (2017): Ch 14.5

USArrests Data



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Principal Component Analysis (PCA)

$$Z_1 = \phi_{11}X_1 + \phi_{21}X_2 + \dots + \phi_{p1}X_p$$
$$z_{i1} = \phi_{11}x_{i1} + \phi_{21}x_{i2} + \dots + \phi_{p1}x_{ip}$$

$$\max_{\phi_{11},...,\phi_{p1}} \{ \frac{1}{n} \sum_{i=1}^{n} (\sum_{j=1}^{p} \phi_{j1} x_{ij})^{2} \}$$

subject to
$$\sum\limits_{j=1}^p \phi_{j1}^2 = 1$$

$$z_{i2} = \phi_{12}x_{i1} + \phi_{22}x_{i2} + \dots + \phi_{p2}x_{ip}$$

Singular Value Decomposition (SVD)

$$X_{n imes p} = U_{n imes p} D_{p imes p} V_{p imes p}^T$$
 U and V are Orthogonal
 $U^T U = I_{n imes n}$ and $V^T V = I_{p imes p}$
 $S = X^T X = V D^2 V^T$
 $X X^T = U D^2 U^T$

 $(S - \delta I)v = 0$

$$z_1 = Xv_1 = u_1d_1$$

 $Var(z_1) = \frac{d_1^2}{n}$

Subsequent Principal Components z_j have maximum variance $\frac{d_j^2}{n}$, subject to being orthogonal to the earlier ones

OLS and Ridge Fitted Vector

$$X\hat{\beta}^{ls} = X(X^TX)^{-1}X^Ty$$

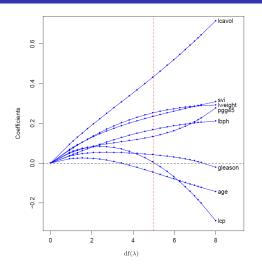
$$= UU^Ty$$

$$X\hat{\beta}^{ridge} = X(X^TX + \lambda I)^{-1}X^Ty$$

$$= UD(D^2 + \lambda I)^{-1}DU^Ty$$

$$= \sum_{j=1}^{p} u_j \frac{d_j^2}{d_j^2 + \lambda} u_j^Ty$$

$$df(\lambda) = \sum_{j=1}^{p} \frac{d_j^2}{d_j^2 + \lambda} = tr[X(X^TX + \lambda I)^{-1}X^T]$$



Effective Degrees of Freedom

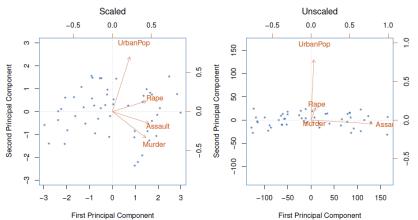
First and Second Principal Component

| | PC1 | PC2 |
|----------|-----------|------------|
| Murder | 0.5358995 | -0.4181809 |
| Assault | 0.5831836 | -0.1879856 |
| UrbanPop | 0.2781909 | 0.8728062 |
| Rape | 0.5434321 | 0.1673186 |

Scaling the Variables

Assault per 100 people rather per 100,00 people

Variance for Murder, Rape, Assault, and UrbanPop: 18.97, 87.73, 6945.16, and 209.5



Proportion of Variance Explained (PVE)

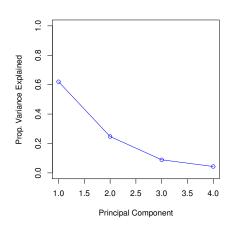
$$PVE = \frac{\frac{\frac{1}{n}\sum\limits_{i=1}^{n}z_{im}^{2}}{\sum\limits_{j=1}^{p}Var(X_{j})}$$

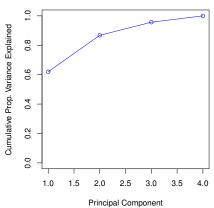
$$\sum_{j=1}^{p} Var(X_{j}) = \sum_{j=1}^{p} \frac{1}{n} \sum_{i=1}^{n} x_{ij}^{2}$$

$$\frac{1}{n}\sum_{i=1}^{n}z_{im}^{2}=\frac{1}{n}\sum_{i=1}^{n}\left(\sum_{j=1}^{p}\phi_{jm}x_{ij}\right)^{2}$$

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Cumulative Proportion of Variance Explained





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Means and Standard Deviations

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

pr.out\$rotation=-pr.out\$rotation

pr.out\$x=-pr.out\$x
pr.out\$rotation

```
        PC1
        PC2
        PC3
        PC4

        Murder
        0.5358995
        -0.4181809
        0.3412327
        -0.64922780

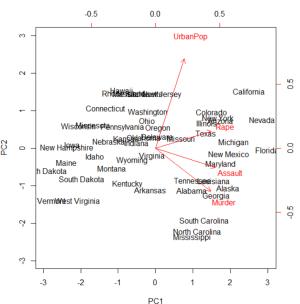
        Assault
        0.5831836
        -0.1879856
        0.2681484
        0.74340748

        UrbanPop
        0.2781909
        0.8728062
        0.3780158
        -0.1338773

        Rape
        0.5434321
        0.1673186
        -0.8177779
        -0.08902432
```

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biplot(pr.out, scale=0)



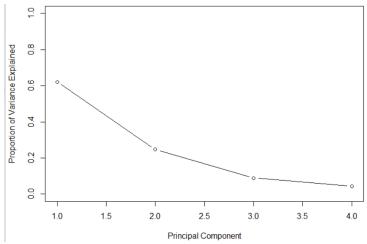
pr.var=pr.out\$sdev^2; pr.var

2.4802416 0.9897652 0.3565632 0.1734301

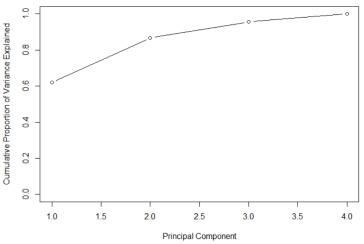
pve=pr.var/sum(pr.var); pve

0.62006039 0.24744129 0.08914080 0.04335752

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')



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6,830 Gene Expression on 64 Cancer Cell Lines

library(ISLR)
nci.labs=NCl60\$labs; nci.data=NCl60\$data
dim(nci.data)

64 6830

table(nci.labs)

```
        BREAST
        CNS
        COLON K562A-repro
        K562B-repro

        7
        5
        7
        1
        1

        LEUKEMIA MCF7A-repro MCF7D-repro MCFAD-repro
        MELANOMA MELANOMA
```

pr.out=prcomp(nci.data, scale=TRUE)

summary (pr.out)

```
Importance of components:
```

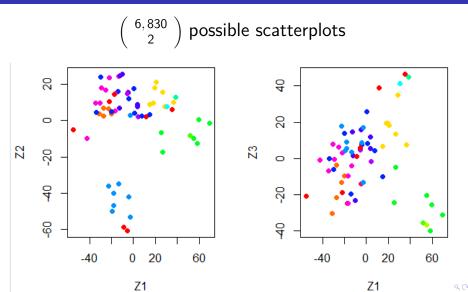
```
PC1
                                      PC2
                                               PC3
                                                         PC4
Standard deviation
                        27.8535 21.48136 19.82046 17.03256
Proportion of Variance
                         0.1136
                                 0.06756
                                           0.05752
                                                    0.04248
Cumulative Proportion
                                 0.18115
                                           0.23867
                         0.1136
                                                    0.28115
                             PC5
                                       PC6
                                                PC7
                                                          PC8
Standard deviation
                        15.97181 15.72108 14.47145 13.54427
Proportion of Variance
                         0.03735
                                  0.03619
                                            0.03066
                                                     0.02686
Cumulative Proportion
                                  0.35468
                                            0.38534
                         0.31850
                                                     0.41220
                             PC9
                                      PC10
                                               PC11
                                                         PC12
Standard deviation
                        13.14400 12.73860 12.68672 12.15769
Proportion of Variance
                         0.02529
                                  0.02376
                                            0.02357
                                                     0.02164
Cumulative Proportion
                         0.43750
                                  0.46126
                                            0.48482
                                                     0.50646
```

Assign a color to each of the 64 cell lines

```
Cols=function (vec ){
 cols=rainbow (length (unique (vec )))
 return (cols[as.numeric (as.factor (vec))])
par(mfrow = c(1,2))
plot(pr.out$x [,1:2], col = Cols(nci.labs), pch = 19,
 xlab = "Z1", ylab = "Z2")
plot(pr.outx[,c(1,3)], col = Cols(nci.labs),
 pch =19, xlab ="Z1",ylab="Z3").....
```

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Projections of the NCI60 cancer cell lines onto the first three principal components



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