12 - PCA Additional Example

Junvie Pailden

SIUE, F2017, Stat 589

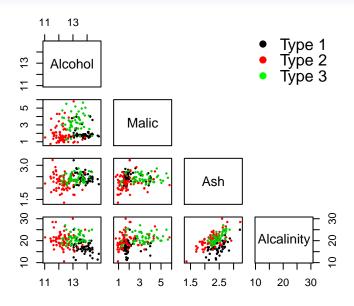
October 02, 2017

Wine Data

The wine data is the result of a chemical analysis of wines grown in the same region in Italy but derived from three different cultivars. The analysis determined the quantities of 13 constituents found in each of the three types of wines.

```
data(wine, package = 'rattle.data')
names(wine)
```

```
[1]
       "Type"
                          "Alcohol"
                                              "Malic"
   Γ41
#
       "Ash"
                          "Alcalinity"
                                              "Magnesium"
   [7] "Phenols"
                          "Flavanoids"
                                              "Nonflavanoids"
                                              "Hue"
  [10] "Proanthocyanins" "Color"
  [13] "Dilution"
                          "Proline"
```



PCA Wine Data using prcomp

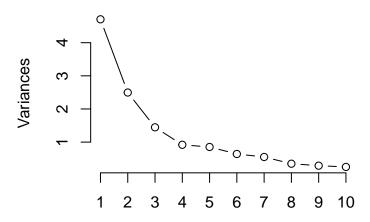
```
# need to center and scale the data
wine.pca1 <- prcomp(wine[,-1], scale. = T, center = T)
names(wine.pca1)

# [1] "sdev" "rotation" "center" "scale" "x"

# Variances/Eigenvalues of PCA
(wine.pca1$sdev)^2</pre>
```

[1] 4.71 2.50 1.45 0.92 0.85 0.64 0.55 0.35 0.29 0.25 0

wine.pca1



Summary of PCA

```
summary(wine.pca1)
```

Importance of components%s:

```
PC2
#
                           PC1
                                       PC3
                                              PC4
                                                     PC5
                         2.169 1.580 1.203 0.9586 0.9237 0
 Standard deviation
 Proportion of Variance 0.362 0.192 0.111 0.0707 0.0656 0
 Cumulative Proportion
                         0.362 0.554 0.665 0.7360 0.8016 0
#
                            PC8
                                   PC9
                                         PC10
                                                PC11
 Standard deviation
                         0.5903 0.5375 0.5009 0.4752 0.411
 Proportion of Variance 0.0268 0.0222 0.0193 0.0174 0.013
 Cumulative Proportion
                         0.9202 0.9424 0.9617 0.9791 0.992
```

First Four Eigenvectors (or Loadings) using prcomp

wine.pca1\$rotation[,1:4]

```
#
                     PC1
                             PC2
                                   PC3
                                          PC4
 Alcohol
                 -0.1443
                          0.4837 - 0.207
                                        0.018
 Malic
                  0.2452
                          0.2249 0.089 -0.537
# Ash
                  0.0021
                          0.3161 0.626 0.214
 Alcalinity
                  0.2393 -0.0106 0.612 -0.061
 Magnesium
               -0.1420
                          0.2996 0.131 0.352
# Phenols
               -0.3947
                          0.0650 0.146 -0.198
 Flavanoids -0.4229 -0.0034 0.151 -0.152
# Nonflavanoids
                  0.2985
                          0.0288
                                 0.170 0.203
 Proanthocyanins -0.3134 0.0393 0.149 -0.399
# Color
                  0.0886
                          0.5300 - 0.137 - 0.066
                 -0.2967 -0.2792 0.085 0.428
 Hue
 Dilution
                 -0.3762 -0.1645 0.166 -0.184
 Proline
                 -0.2868
                          0.3649 - 0.127
                                        0.232
```

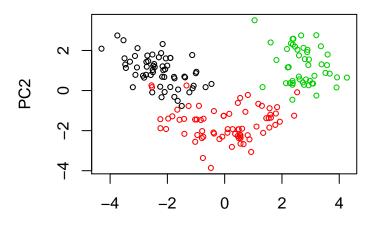
First Four Principal Components (or Scores) using prcomp

```
head(wine.pca1$x[,1:4], 10)
```

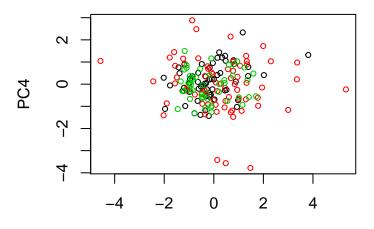
```
#
        PC1 PC2 PC3 PC4
#
  [1.] -3.3 1.44 -0.17 0.215
#
   [2,] -2.2 -0.33 -2.02 0.291
#
  [3,] -2.5 1.03 0.98 -0.723
# [4,] -3.7 2.75 -0.18 -0.566
# [5,] -1.0 0.87 2.02 0.409
# [6,] -3.0 2.12 -0.63 0.514
# [7,] -2.4 1.17 -0.97 0.066
# [8,] -2.1 1.60 0.15 1.189
# [9,] -2.5 0.92 -1.77 -0.056
 [10.] -2.7 0.79 -0.98 -0.348
```

plot(wine.pca1\$x[,1:2], col = wine\$Type, cex = 0.7, main =

1st and 2nd PC's



3rd and 4th PC's



PCA Wine Data using princomp

```
# set cor = TRUE to use scaled data
wine.pca2 <- princomp(wine[,-1], cor = TRUE)
names(wine.pca2)
# [1] "sdev"
                 "loadings" "center" "scale" "n.obs"
# [7] "call"
# Variances/Eigenvalues of PCA
(wine.pca2$sdev)^2
```

```
# Comp.1 Comp.2 Comp.3 Comp.4 Comp.5 Comp.6 Comp.7
# 4.71 2.50 1.45 0.92 0.85 0.64 0.55
# Comp.10 Comp.11 Comp.12 Comp.13
# 0.25 0.23 0.17 0.10
```

First Four Eigenvectors (or Loadings) using princomp

wine.pca2\$loadings[,1:4]

```
#
                 Comp.1 Comp.2 Comp.3 Comp.4
# Alcohol
                -0.1443 -0.4837 -0.207 -0.018
# Malic
                 0.2452 -0.2249 0.089 0.537
# Ash
                 0.0021 -0.3161 0.626 -0.214
 Alcalinity 0.2393 0.0106 0.612 0.061
           -0.1420 -0.2996 0.131 -0.352
 Magnesium
# Phenols
         -0.3947 -0.0650 0.146 0.198
# Flavanoids -0.4229 0.0034 0.151 0.152
# Nonflavanoids 0.2985 -0.0288 0.170 -0.203
 Proanthocyanins -0.3134 -0.0393 0.149 0.399
# Color
                 0.0886 -0.5300 -0.137 0.066
                -0.2967 0.2792 0.085 -0.428
# Hue
 Dilution
                -0.3762 0.1645 0.166 0.184
# Proline
               -0.2868 -0.3649 -0.127 -0.232
```

First Four Principal Components (or Scores) using princomp

head(wine.pca2\$scores[,1:4], 10)

```
#
       Comp.1 Comp.2 Comp.3 Comp.4
#
  [1,]
        -3.3 -1.44 -0.17 -0.216
  [2,] -2.2 0.33 -2.03 -0.291
#
#
  [3,] -2.5 -1.03 0.98 0.725
#
  [4,] -3.8 -2.76 -0.18 0.568
#
  [5.] -1.0 -0.87 2.03 -0.410
  [6.] -3.1 -2.12 -0.63 -0.516
#
#
  [7.] -2.4 -1.17 -0.98 -0.066
  [8.] -2.1 -1.61 0.15 -1.193
#
  [9.] -2.5 -0.92 -1.77 0.056
#
 [10.] -2.8 -0.79 -0.98 0.349
```