

dimensional reduction

dimensional reduction

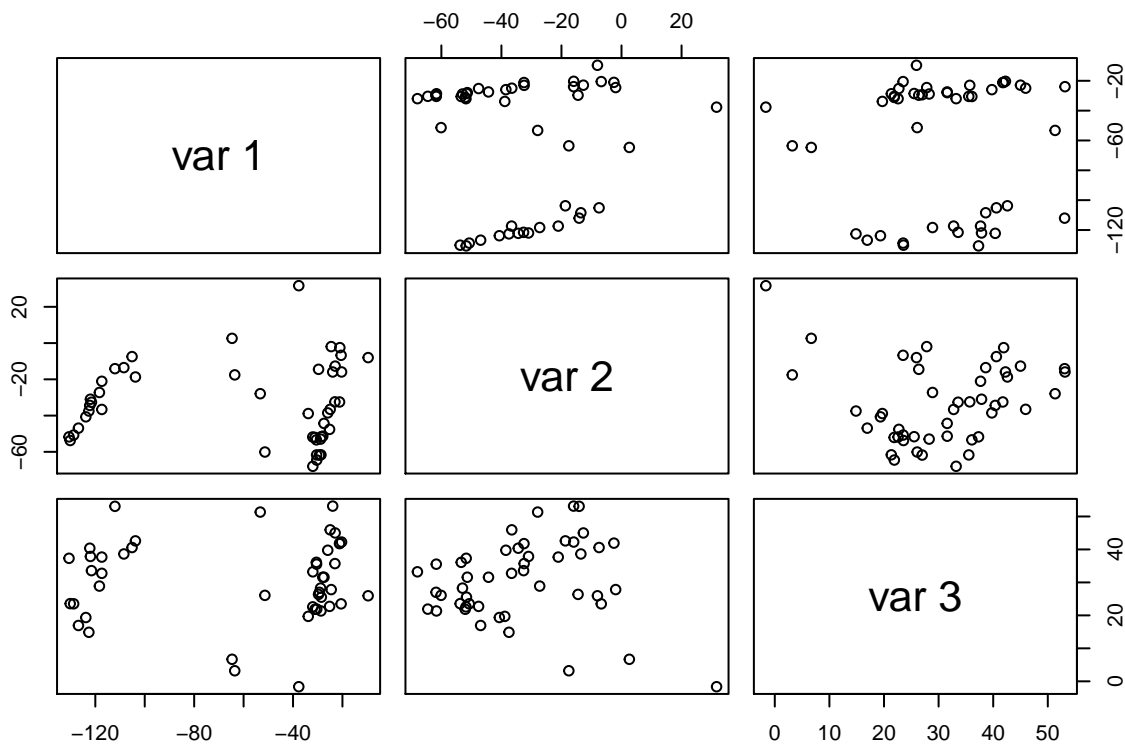
principle component analysis (PCA)

let's first try to code PCA:

```
myPCA <- function(x, d=2){  
  #step 1: covariance estimation  
  #scale function to center the data: scale=False, not normalize, only centered  
  xbar=scale(x,center=TRUE,scale = FALSE)  
  sigma=1/nrow(xbar)*t(xbar)%*%xbar  
  
  #step 2: eigen-decomposition  
  out=eigen(sigma)  
  
  #step 3: return the PC axes  
  return(out$vector[,1:d])  
  #return(out)  
}
```

now let's try it on some data:

```
data("swiss")  
u=myPCA(swiss,d=3)  
  
  #now we have to project the data  
  xproj=as.matrix(swiss) %*% u  
  #xproj  
  #plot(xproj, type='p', pch=19) # for 1-2 dimension  
  pairs(xproj)#for more than 2 dimension
```



#so we move from 6 dimensional spaces to other number of dimensional spaces

now we can combine the plotting into the function

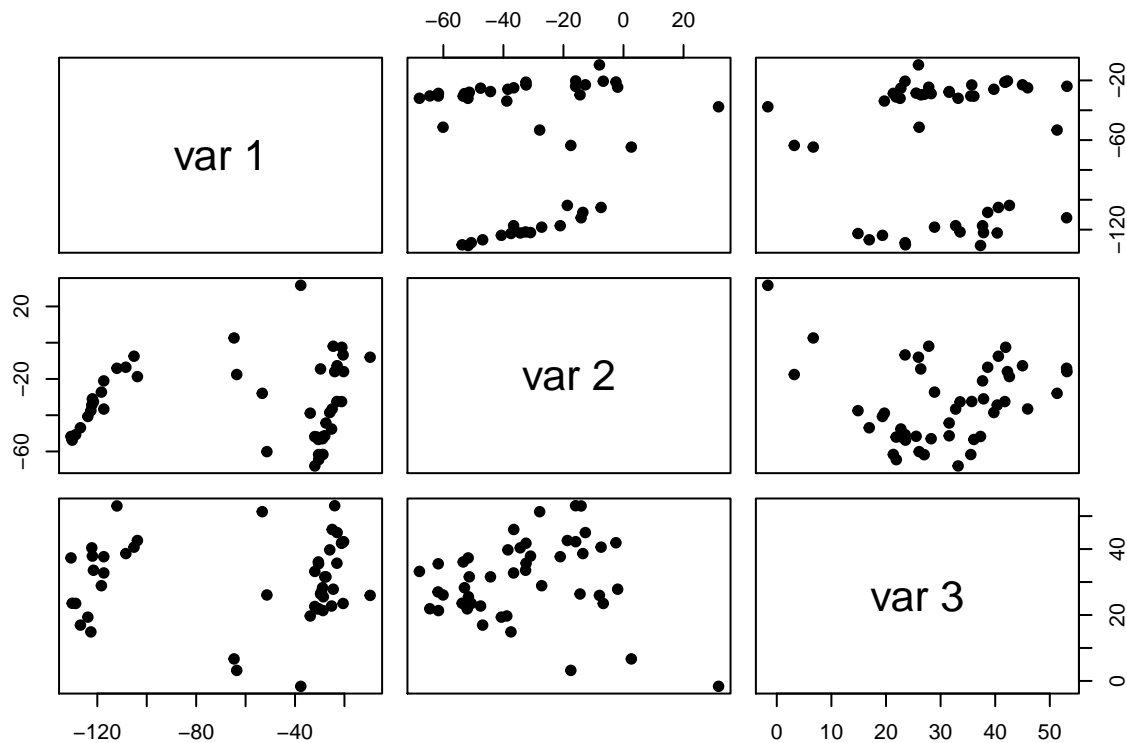
```
myPCAPlot <- function(x, d=2){
  #step 1: covariance estimation
  #scale function to center the data: scale=False, not normalize, only centered
  xbar=scale(x,center=TRUE,scale = FALSE)
  sigma=1/nrow(xbar)*t(xbar)%*%xbar

  #step 2: eigen-decomposition
  out=eigen(sigma)

  #step 3: return the PC axes
  #return(out$vector[,1:d])
  #return(out)
  u=out$vector[,1:d]
  xproj=as.matrix(x) %*% u
  if(d<=2)
    plot(xproj, type = 'p', pch = 19)
  else
    pairs(xproj,pch=19)
  return(list(u=u,lambda=out$values))
}
```

try with swiss

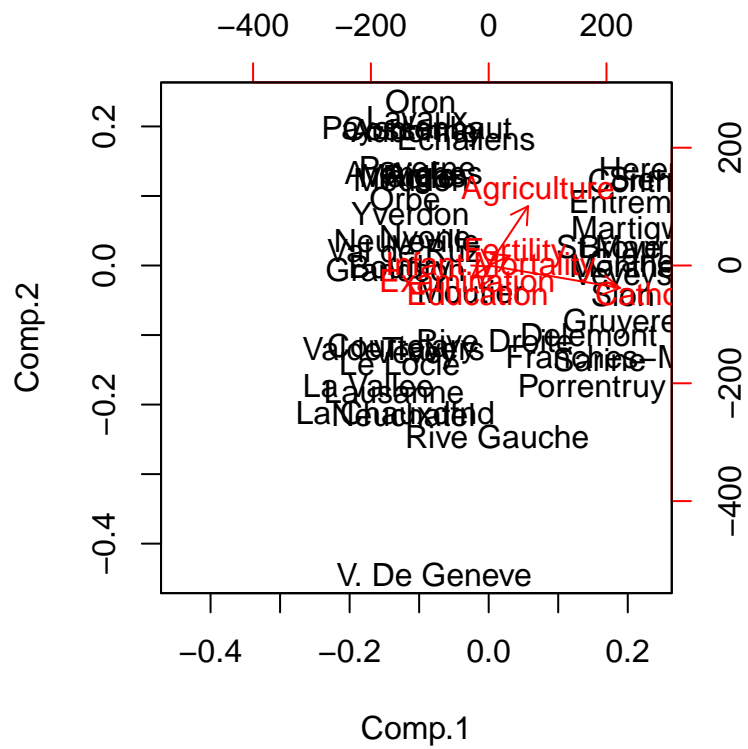
```
myPCAPlot(swiss,d=3)
```



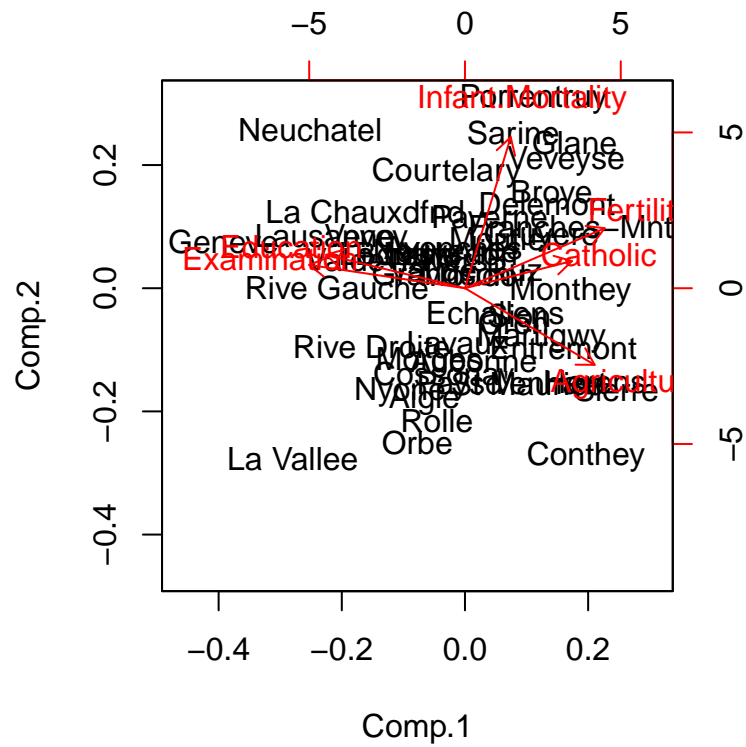
```
## $u
##           [,1]      [,2]      [,3]
## [1,] -0.15163143 -0.14270789  0.81454413
## [2,] -0.28121756 -0.85914886 -0.35256541
## [3,]  0.12207834  0.17688621 -0.18767793
## [4,]  0.06329733  0.32260928 -0.40096045
## [5,] -0.93748965  0.32543441 -0.07870742
## [6,] -0.01131739  0.01498883  0.10014161
##
## $lambda
## [1] 1880.67818  456.72826  142.19366   22.16770   13.09300    6.05952
```

if we use the R function 'princomp':

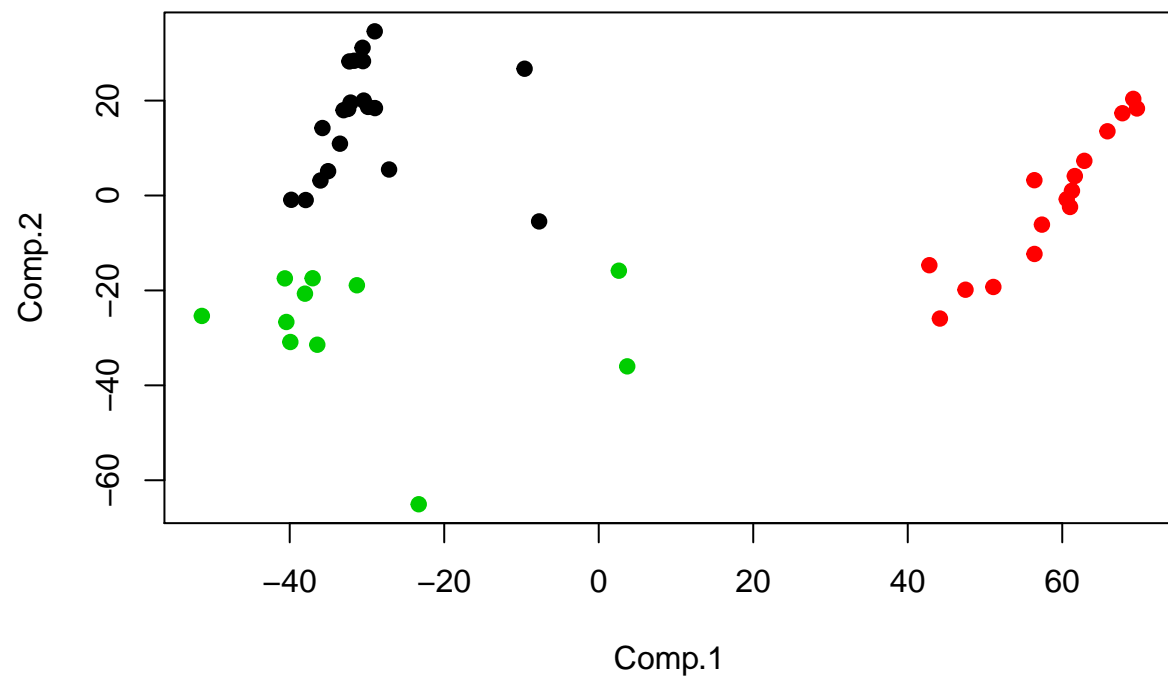
```
#comparision between PCA and scaled PCA:
out=princomp(swiss)
out1=princomp(swiss,cor = TRUE)#scaleD ??????????
xproj=predict(out,swiss)
#par(mfrow=c(1,2))
biplot(out)
```



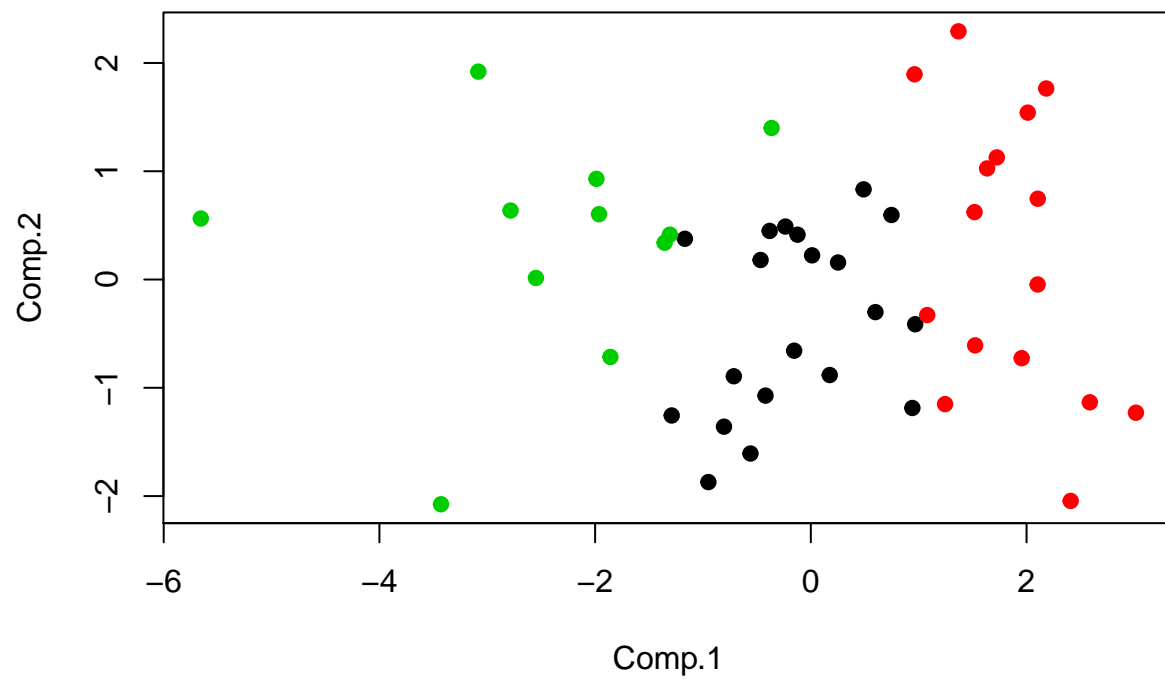
```
biplot(out1)
```



```
clus=kmeans(swiss,3)
plot(predict(out,swiss),col=clus$cluster,pch=19)
```



```
plot(predict(out1,swiss),col=clus$cluster,pch=19) #once it's scaled, the cluster is not that obvious
```

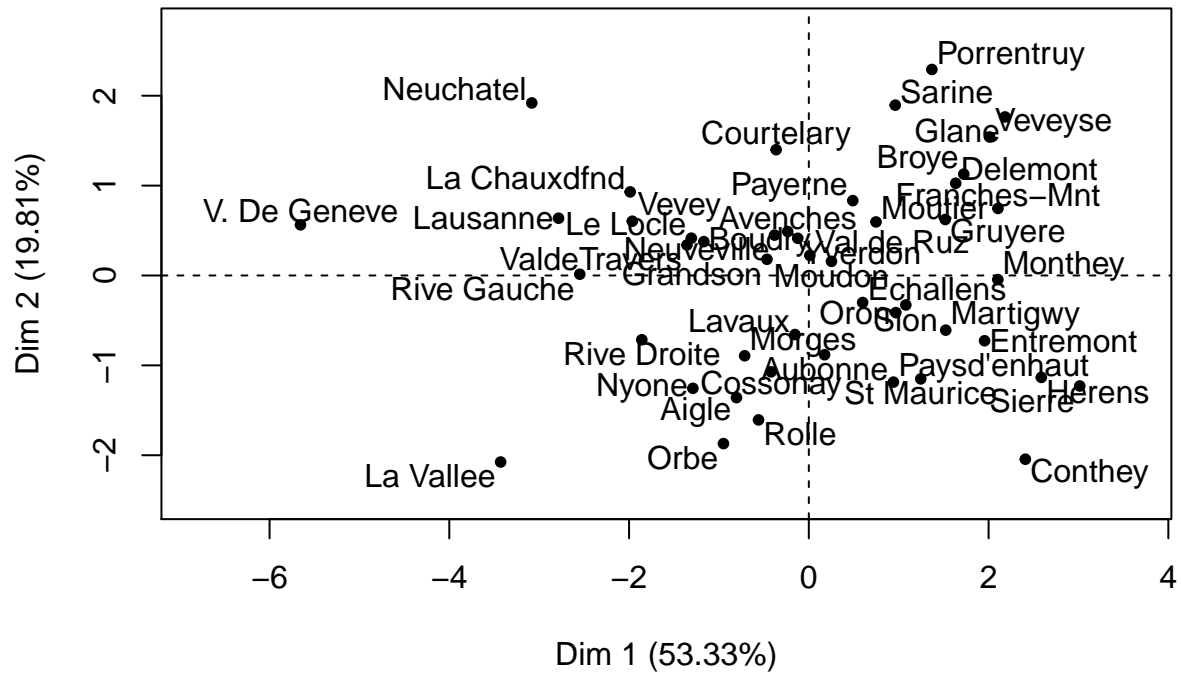


#it is not clear since it's combining the 2 plot

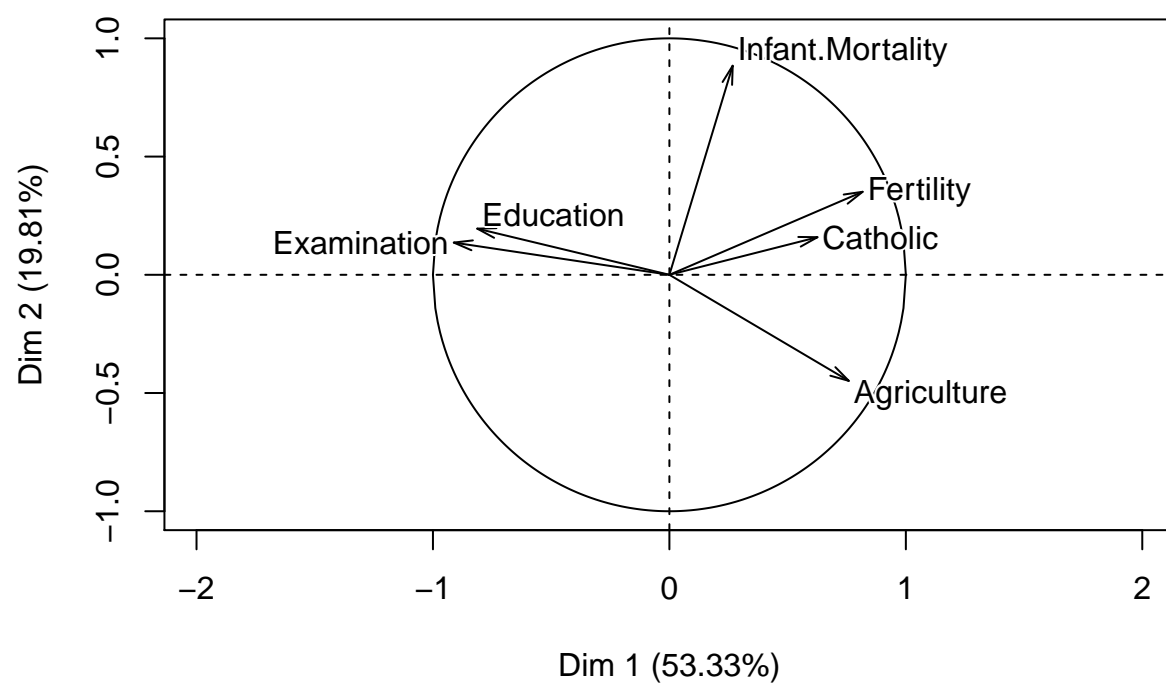
in the 'FactoMineR' package several additional visualizations are possible:

```
#install.packages('FactoMineR')
#here unlike biplot it seperate 2 graph
library(FactoMineR)
out=PCA(swiss)
```

Individuals factor map (PCA)

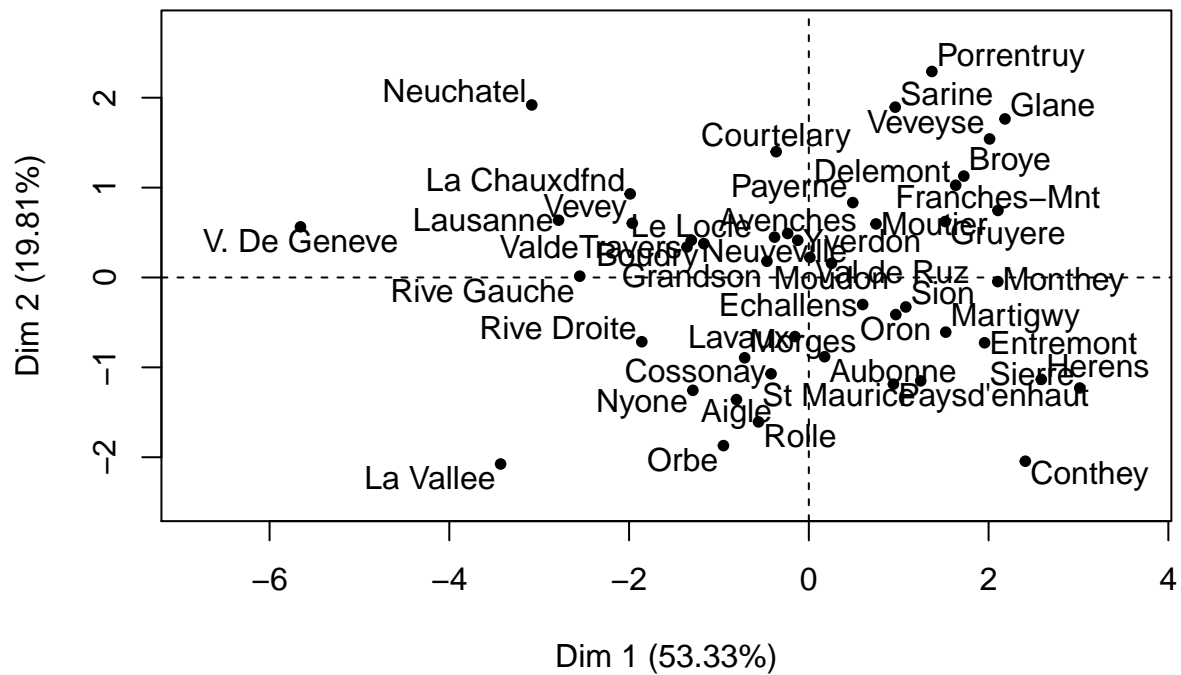


Variables factor map (PCA)



```
plot(out)
```

Individuals factor map (PCA)



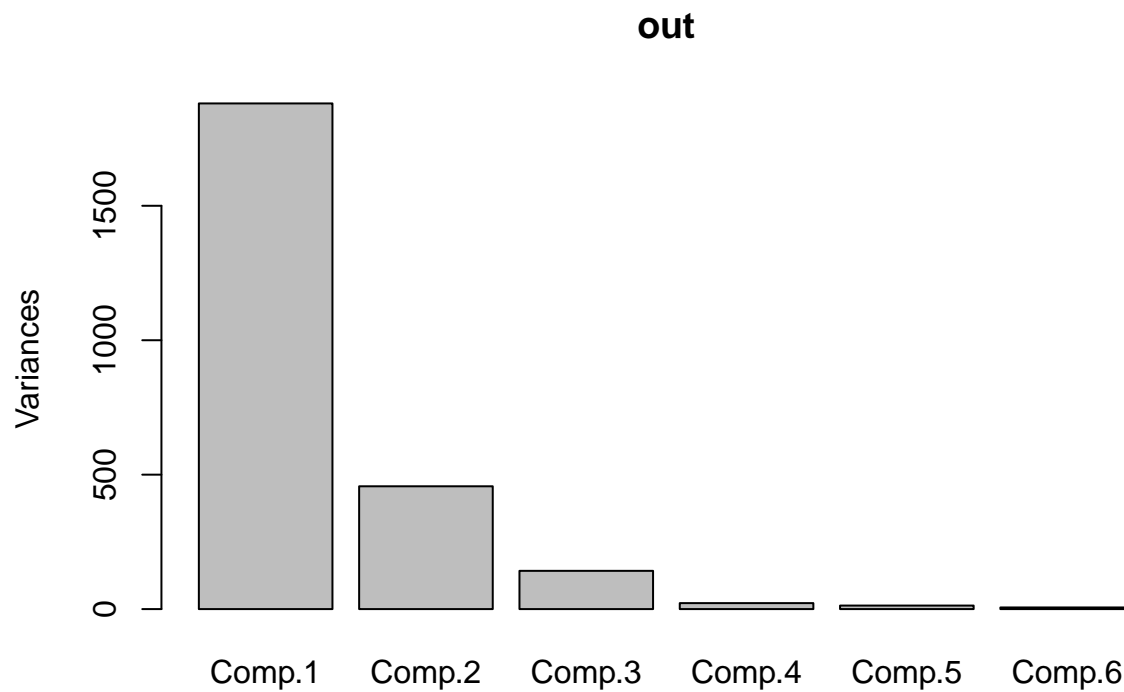
exercise: use the three methods we saw in class for choosing the right number of components to retain

90% rule, eigenvalue scree:

```
out=princomp(swiss)
out$sdev
```

```
##      Comp.1      Comp.2      Comp.3      Comp.4      Comp.5      Comp.6
## 43.366787 21.371202 11.924498  4.708259  3.618425  2.461609
```

```
plot(out) #variance of the variable
```



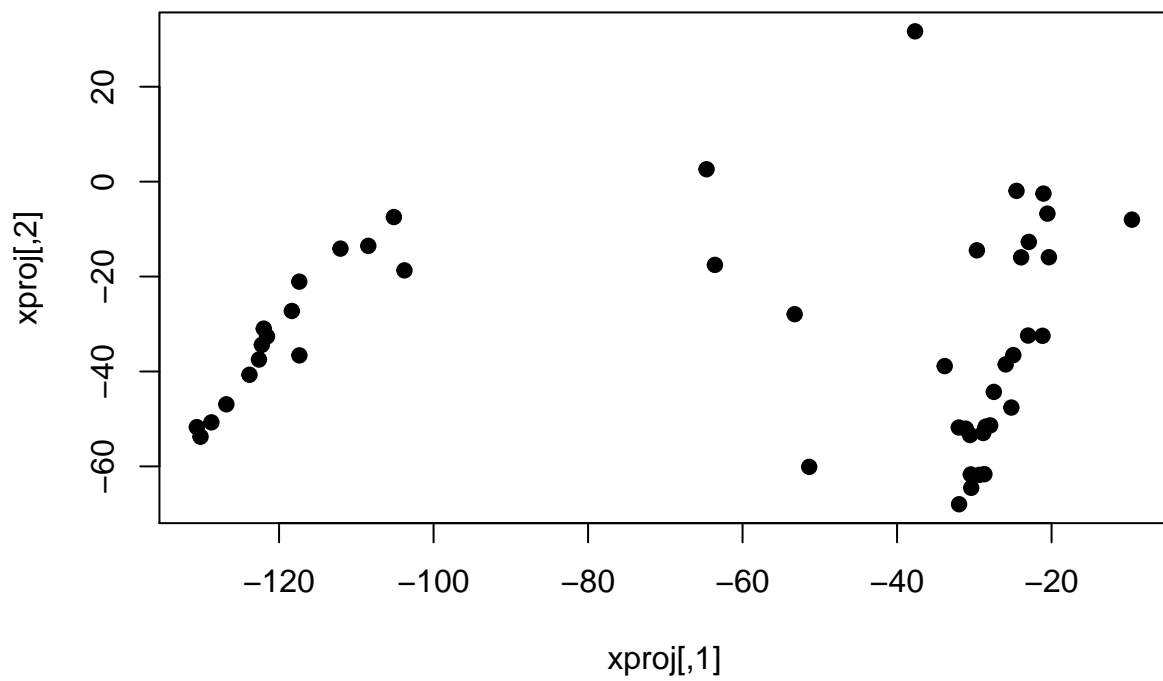
```
summary(out)
```

```
## Importance of components:
##               Comp.1    Comp.2    Comp.3    Comp.4
## Standard deviation 43.3667866 21.3712016 11.92449843 4.708258751
## Proportion of Variance 0.7460284 0.1811752 0.05640546 0.008793495
## Cumulative Proportion 0.7460284 0.9272036 0.98360907 0.992402568
##               Comp.5    Comp.6
## Standard deviation 3.618425207 2.461609258
## Proportion of Variance 0.005193739 0.002403694
## Cumulative Proportion 0.997596306 1.000000000
```

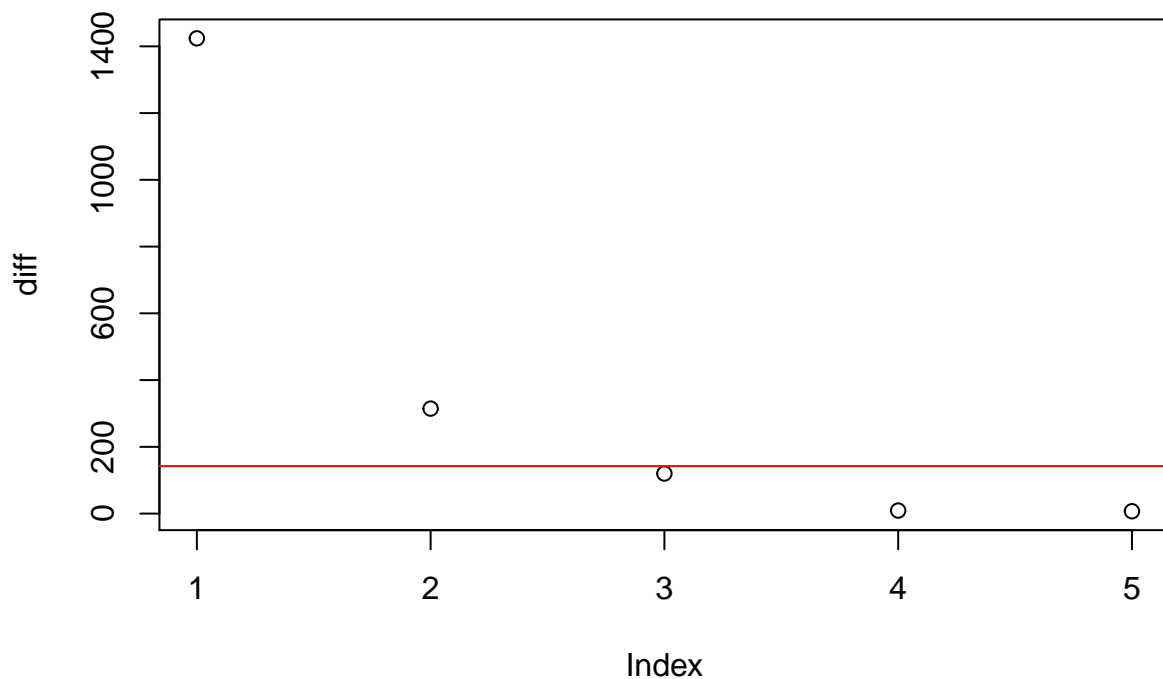
```
# with the 90% rule we can take already first 2 variable
#eigenvalue scree: 3 is still important so choose between 2-3
```

The Cattell's test:

```
out=myPCAPlot(swiss)
```



```
diff=abs(diff(out$lambda))
plot(diff)
abline(h=0.1*max(diff),col='red')
```



#cattell say we should retain 2+1=3 dimension

MDS: Multi dimensional scaling

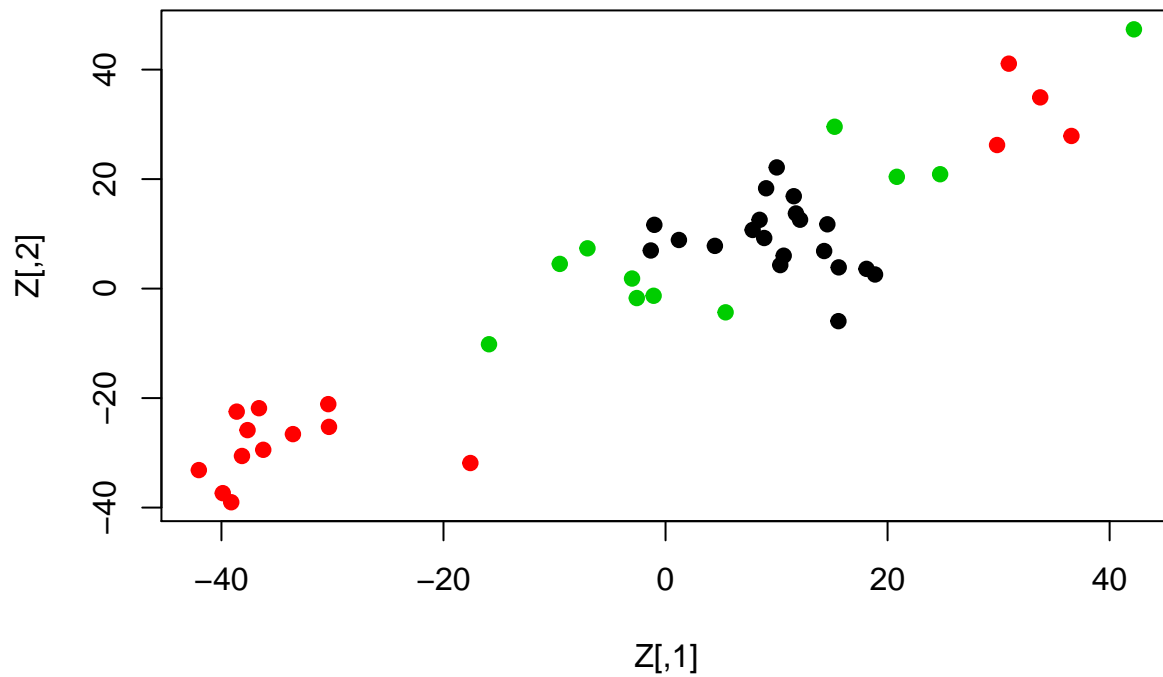
```
myMDS<- function(x,d=2){
  n=nrow(x);
  D =as.matrix(dist(x)) #dist() for continuous data, if it's other type, we need to find the appropriate
  fun<-function(par,D){
    Z=matrix(par,ncol=d)
    s=0
    for(i in 1:n)
      for(j in 1:n)
        s=s+(D[i,j]-sqrt(sum(Z[i,]-Z[j,])^2))^2
    return(s) #return s
  }
  zstart=matrix(runif(n*d),ncol=d)
  out=optim(zstart,fun,D=D,method='SANN')
  Z=out$par
}

Z=myMDS(swiss,d=2)
Z
```

```
##           [,1]      [,2]
```

```
## [1,] -1.072692 -1.311884
## [2,] 29.862481 26.223676
## [3,] -30.311328 -25.262600
## [4,] 10.016776 22.131668
## [5,] 4.446070 7.800568
## [6,] -17.579059 -31.863214
## [7,] -38.623252 -22.475850
## [8,] -36.234708 -29.448289
## [9,] -36.619207 -21.847400
## [10,] -30.376528 -21.108971
## [11,] 36.559829 27.880655
## [12,] 18.871348 2.573557
## [13,] 12.115318 12.580564
## [14,] 8.471652 12.559274
## [15,] 14.576332 11.739828
## [16,] -1.326661 6.960145
## [17,] 15.573996 -5.946418
## [18,] -2.588884 -1.732843
## [19,] -9.525880 4.508308
## [20,] 9.061513 18.322361
## [21,] 14.286759 6.860202
## [22,] 7.848856 10.709643
## [23,] 10.334114 4.290972
## [24,] 15.599655 3.875896
## [25,] 11.551308 16.871476
## [26,] 8.889130 9.240755
## [27,] 11.747680 13.722272
## [28,] 18.103042 3.605089
## [29,] 5.411410 -4.336935
## [30,] 10.643379 6.008368
## [31,] -42.035979 -33.150837
## [32,] 30.921202 41.082104
## [33,] -39.880415 -37.368111
## [34,] -38.150340 -30.575192
## [35,] -37.650347 -25.851444
## [36,] 33.750006 34.926526
## [37,] -39.117722 -39.025319
## [38,] -33.560205 -26.582765
## [39,] -1.006708 11.645277
## [40,] 15.230159 29.560467
## [41,] -3.015084 1.832691
## [42,] 24.733879 20.891635
## [43,] 1.207141 8.879745
## [44,] -7.030778 7.351308
## [45,] 42.191971 47.351505
## [46,] 20.834148 20.418903
## [47,] -15.900820 -10.168220
```

```
plot(Z,type='p',pch=19,col=clus$cluster)
```



#MDS is very time consuming

- MDS in R 'cmdscale' Much better in performance

```
d=dist(swiss)
res=cmdscale(d,k=2,eig=TRUE)
res
```

```
## $points
##           [,1]      [,2]
## Courtelary   37.032433 -17.4348788
## Delemont    -42.797334 -14.6876683
## Franches-Mnt -51.081639 -19.2740356
## Moutier       7.716707  -5.4587215
## Neuveville   35.032658   5.1260970
## Porrentruy  -44.161953 -25.9224124
## Broye        -56.392984   3.2255060
## Glane        -61.258244   0.9998919
## Gruyere      -56.405711 -12.3159788
## Sarine       -47.477237 -19.8509107
## Veveyse      -61.008008  -2.4123170
## Aigle        28.965873  18.4219674
## Aubonne      31.653458  28.3931122
## Avenches     32.123633  19.5819402
## Cossonay     32.274690  28.2546346
```

```

## Echallens      9.595877  26.7118107
## Grandson      39.800148  -0.9053319
## Lausanne      40.435132 -26.6464679
## La Vallee     51.376323 -25.3760191
## Lavaux       30.572130  31.1329195
## Morges       32.435116  18.2481593
## Moudon       33.028767  17.9772680
## Nyone        27.138257   5.4808462
## Orbe         35.765096  14.2149538
## Oron        28.996500  34.5897700
## Payerne      30.425130  20.0025379
## Paysd'enhaut  30.515769  28.3119655
## Rolle       29.864476  18.6702353
## Vevey       31.307144 -18.9143568
## Yverdon     33.499133  10.9174996
## Conthey     -67.788045  17.3371596
## Entremont   -65.850277  13.5345554
## Herens      -69.193539  20.3626868
## Martigwy    -62.858166   7.3053848
## Monthey     -60.583087  -0.7701055
## St Maurice  -61.629377   4.0987185
## Sierre      -69.670618  18.3442505
## Sion        -57.369412  -6.1344239
## Boudry      37.933435  -0.9525147
## La Chauxdfnd 39.920865 -30.8689579
## Le Locle    38.027990 -20.6843295
## Neuchatel   36.433011 -31.4410984
## Val de Ruz   36.020078   3.1624867
## ValdeTravers 40.627884 -17.4614718
## V. De Geneve 23.301944 -65.0529176
## Rive Droite -2.605495 -15.8418221
## Rive Gauche -3.688530 -35.9996173
##
## $eig
## [1]  8.839187e+04  2.146623e+04  6.683102e+03  1.041882e+03  6.153710e+02
## [6]  2.847974e+02  1.718857e-11  1.055757e-11  6.353581e-12  4.313025e-12
## [11] 3.096859e-12  2.205915e-12  2.081644e-12  2.027854e-12  1.995170e-12
## [16] 1.768706e-12  1.710028e-12  1.689229e-12  1.368958e-12  1.292917e-12
## [21] 1.011941e-12  6.688735e-13  6.069093e-13  5.290955e-13  3.429863e-13
## [26] 1.641147e-13  7.494036e-14  3.532351e-14 -8.270456e-14 -3.319275e-13
## [31] -4.524524e-13 -5.877947e-13 -9.702069e-13 -1.068181e-12 -1.106508e-12
## [36] -1.174723e-12 -1.205118e-12 -1.214137e-12 -1.271281e-12 -1.335532e-12
## [41] -1.514998e-12 -1.558048e-12 -1.789720e-12 -1.978327e-12 -2.077321e-12
## [46] -2.263560e-12 -4.255388e-12
##
## $x
## NULL
##
## $ac
## [1] 0
##
## $GOF
## [1] 0.9272036 0.9272036

```



```
plot(res$points,type='p',pch=19,col=clus$cluster)
```

