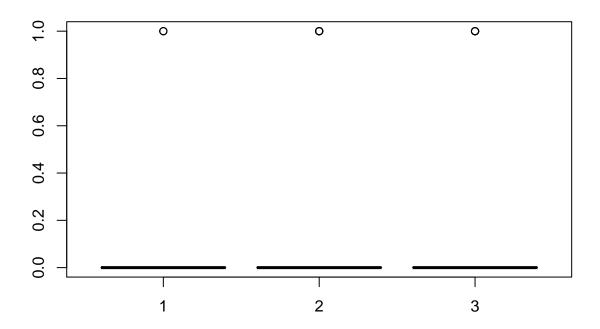
# Supervised learning, Learnin in high dimention

## Supervised learning

#### The general procedure

let's condider some classification problem with the iris data

```
library(MASS); library(class); library(e1071)
x=iris[,-5]
y=iris[,5]
E=matrix(NA,3,nrow(x))
# I would like to compare several classification technique (LDA, KNN, SVM)
#use leave one out method
for (b in 1:nrow(x)){
  xl=x[-b,];yl=y[-b]
  xv=x[b,];yv=y[b] #leave one out
  y1=predict(lda(x1,y1),xv)
  y2=knn(xl,xv,yl)
  #SVM
  y3=predict(svm(x1,y1),xv)
 E[1,b] = as.numeric(y1$class != yv)
 E[2,b] = as.numeric(y2 != yv)
  E[3,b] = as.numeric(y3 != yv)
#pick the most appropriate method
boxplot(t(E))
```



```
## [1] 0.02000000 0.04000000 0.03333333
apply(E,1,sd)
## [1] 0.1404690 0.1966157 0.1801069
#smallest mean and smallest sd, lda is the best method
#my final pick lda
clf=lda(x,y)
\textit{\#in the future when i have new observation}
xstar=c(0.5, 8, 56, 4)
predict(clf,xstar)
## $class
## [1] virginica
## Levels: setosa versicolor virginica
##
## $posterior
     setosa versicolor virginica
## [1,] 0 9.493336e-184
```

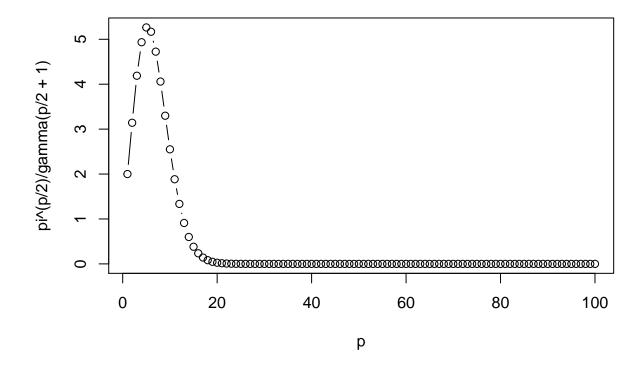
apply(E,1,mean)

### Learning in high-dimensional spaces

#### The curs of dimensionality

The volume of the hyper-sphere is  $V(p) = \frac{\pi^{p/2}}{\Gamma(p/2+1)}$ .

```
p = 1:100
plot(p,pi^(p/2) / gamma(p/2+1), type='b')
```



#### High-dimensional data clustering (HDDC)

HDDC is implemented in the HDclassif library (which also p^roposes the HDDA method for classification).

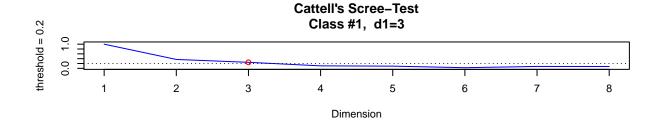
```
#install.packages('HDclassif')
library(HDclassif)
```

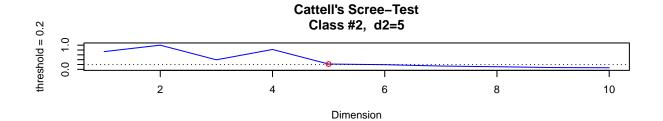
## Warning: package 'HDclassif' was built under R version 3.6.1

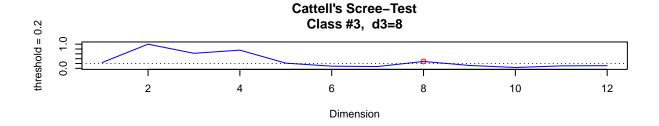
#### #?hddc

We may try HHDC on the wine data set:

```
data(wine)
X = scale(wine[,-1])
Y = wine$class
out = hddc(X,K=1:10)#, threshold = (1:10)/100)
## Warning in hddc_main(model = model, K = K, threshold = threshold, ...):
## Maximum iterations reached (200).
## HDDC:
##
         model K threshold
                                 BIC
## 1 AKJBKQKDK 3 0.2 -5,608.69
## 2 AKJBKQKDK 2
                     0.2 -5,637.93
                     0.2 -5,731.81
## 3 AKJBKQKDK 4
## 4 AKJBKQKDK 5
                     0.2 -5,802.62
## 5 AKJBKQKDK 1
                       0.2 -5,841.07
## 6 AKJBKQKDK 6
                     0.2 -6,005.42
## 7 AKJBKQKDK 7
                     0.2 - 6,175.51
## 8 AKJBKQKDK 8
                     0.2 -6,475.91
## 9 AKJBKQKDK 9
                       0.2 -6,655.57
## 10 AKJBKQKDK 10
                       0.2 -6,776.24
## SELECTED: model AKJBKQKDK with 3 clusters.
## Selection Criterion: BIC.
plot(out)
```







#### out\$d

```
## Intrinsic dimensions of the classes:
## 1 2 3
## dim: 3 5 8
```

#### table(out\$cl,Y)

```
## Y
## 1 2 3
## 1 59 2 0
## 2 0 0 48
## 3 0 69 0
```

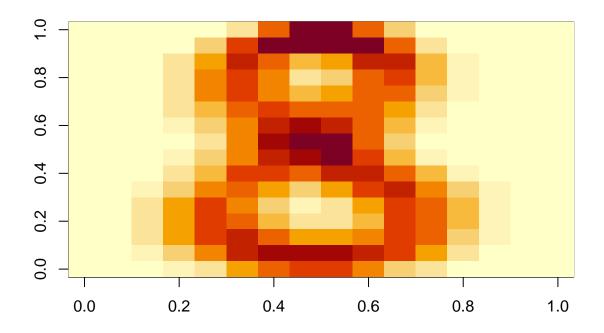
Comparison with Mclust

#### library(mclust)

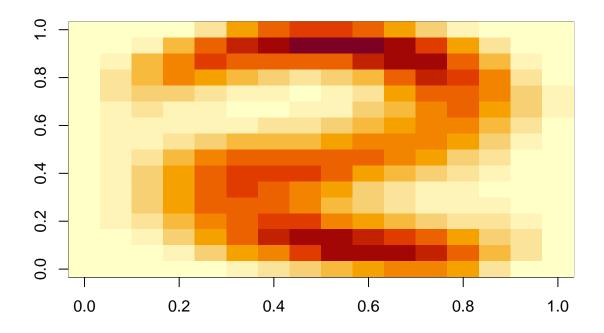
```
## Package 'mclust' version 5.4.3
## Type 'citation("mclust")' for citing this R package in publications.
out1 = hddc(X,K=3)
```

```
model K threshold
## 1 AKJBKQKDK 3 0.2 -5,608.69
## SELECTED: model AKJBKQKDK with 3 clusters.
## Selection Criterion: BIC.
table(out1$cl,Y)
##
     Y
       1 2 3
##
     1 0 69 0
##
    2 0 0 48
   3 59 2 0
##
out2 = Mclust(X,G=3)
table(out2$c1,Y)
##
##
       1 2 3
    1 56 0 0
##
   2 3 70 0
   3 0 1 48
##
Let's now move to higher dimensions:
#install.packages('MBCbook')
library(MBCbook)
## Warning: package 'MBCbook' was built under R version 3.6.1
## Loading required package: Rmixmod
## Loading required package: Rcpp
## Rmixmod v. 2.1.2.2 / URI: www.mixmod.org
## Loading required package: mvtnorm
data("usps358")
X = usps358[,-1]
Y = usps358cls
system.time(out1 <- hddc(X,K=3))</pre>
##
        model K threshold
                                  BIC
## 1 AKJBKQKDK 3 0.2 -613,317.19
## SELECTED: model AKJBKQKDK with 3 clusters.
## Selection Criterion: BIC.
##
     user system elapsed
##
     10.06 0.86 11.30
```

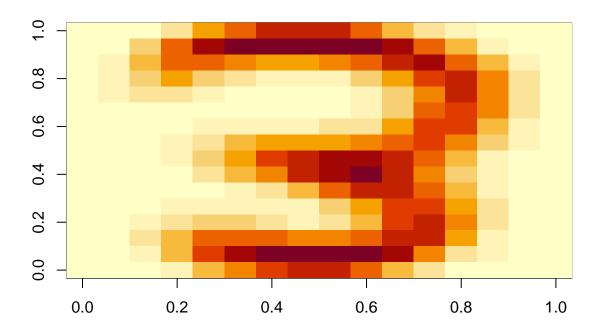
```
table(out1$cl,Y)
##
     Y
            2 3
##
        1
##
    1 596 5 24
##
    2 43 549 26
   3 19 2 492
##
#system.time(out2 <- Mclust(X,G=3,modelNames = 'VVV'))</pre>
#table(out2$cl,Y)
library(MBCbook)
library(HDclassif)
data("usps358")
X = usps358[,-1]
Y = usps358$cls
out1 = hddc(X,K=3)
        model K threshold
## 1 AKJBKQKDK 3 0.2 -613,317.19
## SELECTED: model AKJBKQKDK with 3 clusters.
## Selection Criterion: BIC.
table(out1$cl,Y)
##
##
          2 3
        1
##
   1 19 2 492
    2 43 549 26
##
##
    3 596 5 24
image(matrix(out1$mu[1,],ncol=16))
```



image(matrix(out1\$mu[2,],ncol=16))



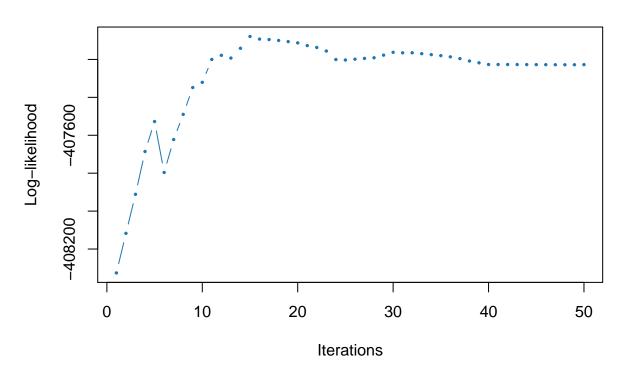
image(matrix(out1\$mu[3,],ncol=16))

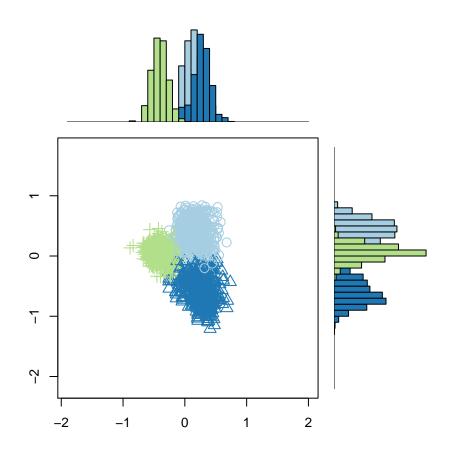


#### $Fisher\ EM$

```
\#install.packages('FisherEM')
library(FisherEM)
## Warning: package 'FisherEM' was built under R version 3.6.1
## Loading required package: parallel
## Loading required package: elasticnet
## Loading required package: lars
## Loaded lars 1.2
out=fem(X,K=3)
table(out$cl,Y)
##
##
                 3
                62
##
##
       48
            34 462
             5 18
##
     3 515
```

# Log-likelihood





## **Group means**

