

DISCRIMINANT ANALYSIS

```

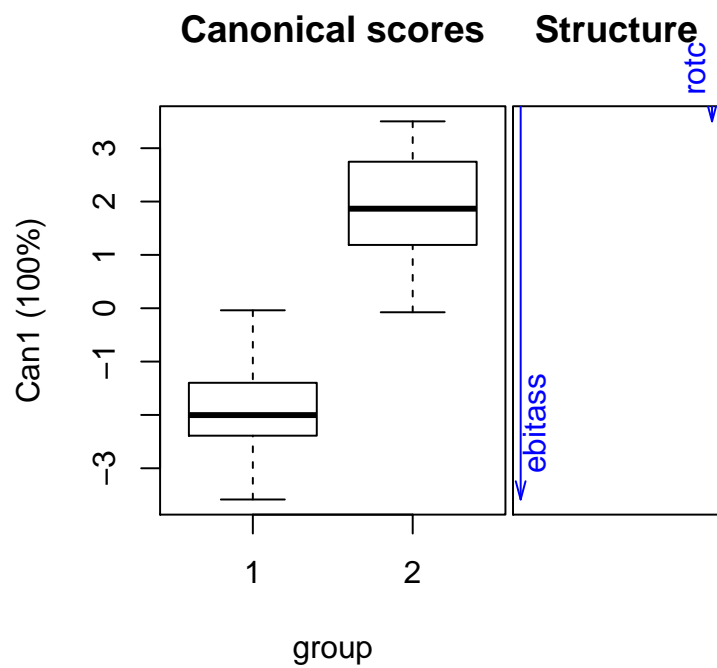
> #####
> #canonical discriminant analysis with 2 groups
> #####
> library(heplots)
> library(candisc)
> ddiscr <- read.table("C:/R/rmmva/data STEPDISCR SHARMA.txt", header=T, quote="\")
> attach(ddiscr)
> ddiscr

  group mktbook  rotc   roe  reass ebitass
1      1  2.304 0.182 0.191 0.377 0.158
2      1  2.703 0.206 0.205 0.469 0.210
3      1  2.385 0.188 0.182 0.581 0.207
4      1  5.981 0.236 0.258 0.491 0.280
5      1  2.762 0.193 0.178 0.587 0.197
6      1  2.984 0.173 0.178 0.546 0.227
7      1  2.070 0.196 0.178 0.443 0.148
8      1  2.762 0.212 0.219 0.472 0.254
9      1  1.345 0.147 0.148 0.297 0.079
10     1  1.716 0.128 0.118 0.597 0.149
11     1  3.000 0.150 0.157 0.530 0.200
12     1  3.006 0.191 0.194 0.575 0.187
13     2  0.975 -0.031 -0.280 0.105 -0.012
14     2  0.945 0.053 0.019 0.306 0.036
15     2  0.270 0.036 0.012 0.269 0.038
16     2  0.739 -0.074 -0.150 0.204 -0.063
17     2  0.833 -0.119 -0.358 0.155 -0.054
18     2  0.716 -0.005 -0.305 0.027 0.000
19     2  0.574 0.039 -0.042 0.268 0.005
20     2  0.800 0.122 0.080 0.339 0.091
21     2  2.028 -0.072 -0.836 -0.185 -0.036
22     2  1.225 0.064 -0.430 -0.057 0.045
23     2  1.502 -0.024 -0.545 -0.050 -0.026
24     2  0.714 0.026 -0.110 0.021 0.016

> cand.mod <- lm(cbind(ebitass, rotc)~ group, data=ddiscr)
> cand.out <- candisc(cand.mod, data=ddiscr)

> plot(cand.out, cex=0.25, cex.lab=0.5)

```



```
> cand.out$eigenvalues
[1] 4.1239e+00 -1.2490e-16

> cand.out$canrsq
[1] 0.80484

> cand.out$coeffs.raw
      Can1
ebitass -15.0919
rotc     -5.7685

> cand.out$coeffs.std
      Can1
ebitass -0.74337
rotc     -0.30547

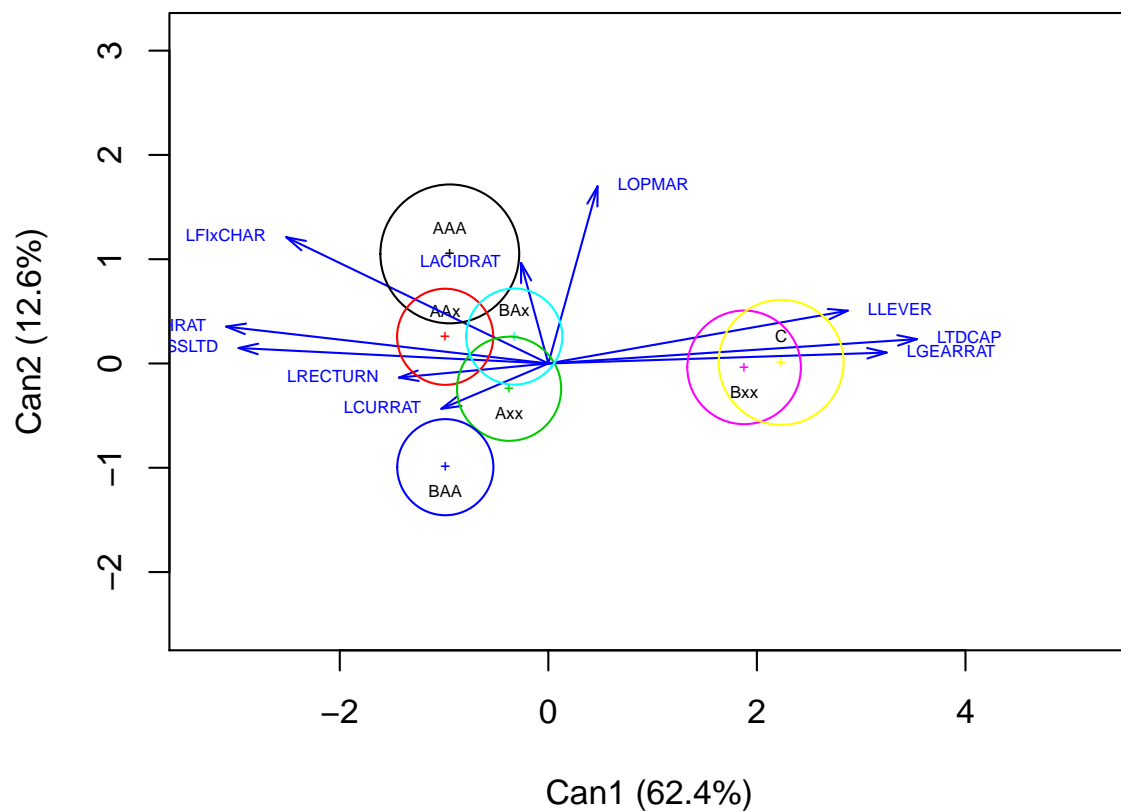
> cand.out$scores[1:5,]
  group  Can1
1     1 -1.4326
2     1 -2.3558
3     1 -2.2067
4     1 -3.5853
5     1 -2.0846
```

```

> #####
> #canonical discriminant analysis with multiple groups
> #####
> training <- read.table("C:/R/rmmva/training data discrim in R.txt", header=T, quote="")
> attach(training)
> cand.mod <- lm(cbind(LOPMAR,LFIxCHAR,LGEARRAT,LTDCAP,LLEVER,LCASHRAT,
+                    LACIDRAT,LCURRAT,RECTURN ,LASSLTD)~ RATING, data=training)
> cand.out <- candisc(cand.mod, data=training)

> plot(cand.out, type="n",scale=4)

```



```

> cand.out$eigenvalues

[1] 1.6912e+00 3.4099e-01 2.6797e-01 1.9603e-01 1.6500e-01 4.9405e-02 9.9297e-15
[8] 1.1861e-15 -1.5733e-16 -1.6638e-15

> cand.out$canrsq

[1] 0.628422 0.254281 0.211338 0.163902 0.141633 0.047079

> cand.out$coeffs.raw

```

	Can1	Can2	Can3	Can4	Can5	Can6
LOPMAR	-0.73584	3.07160	-1.09554	-0.60409	0.64560	1.06205
LFIxCHAR	0.33488	0.93718	2.04368	0.50465	0.49612	-0.44035
LGEARRAT	2.74688	18.34874	3.81957	-33.48078	-4.20339	8.06266
LTD CAP	27.09014	-19.03845	1.52933	24.89427	-13.36769	-22.40470
LLEVER	-5.55828	-7.20063	-4.94711	16.99502	5.30011	-4.16357
LCASHRAT	-0.83967	-3.57219	-0.60746	-0.17672	-2.87384	-2.13627
LACIDRAT	-0.24914	3.41665	-0.69228	0.32935	-0.42113	0.89010
LCURRAT	-2.04495	-2.17102	-1.15283	-0.51191	3.09711	-3.20162
LRECTURN	-1.10688	2.35793	-0.64643	-0.67772	-0.14683	0.88824
LASSLTD	5.88932	18.81681	0.89984	-28.30446	-3.34119	5.16190

```

> #####
> #stepwise selection
> #####
>
> #install.packages("klaR")
> library(klaR)
> training <- read.table("C:/R/rmmva/training data discrim in R.txt", header=T, quote="\")
> attach(training)
> selection <- greedy.wilks(RATING ~ LOPMAR+ LFIxCHAR +LGEARRAT+ LTD CAP
+                               + LLEVER+ LCASHRAT + LACIDRAT +LCURRAT
+                               +LRECTURN +LASSLTD,niveau = 0.20)
> selection

```

Formula containing included variables:

```

RATING ~ LTD CAP + LGEARRAT + LFIxCHAR + LCASHRAT
<environment: 0x000000000947d1e0>

```

Values calculated in each step of the selection procedure:

	vars	Wilks.lambda	F.statistics.overall	p.value.overall	F.statistics.diff
1	LTD CAP	0.50458	12.1095	1.9562e-09	12.1095
2	LGEARRAT	0.42800	6.4305	4.2064e-09	2.1768
3	LFIxCHAR	0.35979	4.9373	3.4038e-09	2.2753
4	LCASHRAT	0.30874	4.1544	3.8332e-09	1.9566
	p.value.diff				
1				1.9562e-09	
2				5.4684e-02	
3				4.5481e-02	
4				8.3235e-02	

```

> #####
> #linear and quadratic classification functions
> #####
> library(MASS)
> ddiscr <- read.table("C:/R/rmmva/data STEPDISCR SHARMA.txt", header=T, quote="\")
> attach(ddiscr)
> #lda uses by default priors proportional to sample sizes
> fitl<- lda(group ~ ebitass + rotc,prior=c(1,1)/2)
> fitl

```

```

Call:
lda(group ~ ebitass + rotc, prior = c(1, 1)/2)

Prior probabilities of groups:
  1  2
0.5 0.5

Group means:
      ebitass      rotc
1 0.1913333 0.18350
2 0.0033333 0.00125

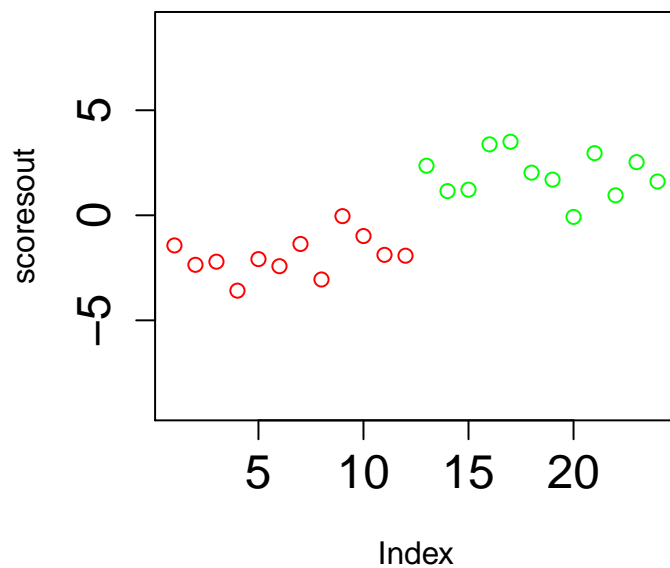
Coefficients of linear discriminants:
      LD1
ebitass -15.0919
rotc     -5.7685

> pp <- predict(fitl,ddiscr)
> table(group,pp$class )

group  1  2
      1 12  0
      2  1 11

> scoresout <-pp$x
> plot(scoresout, col=rainbow(3)[group], asp=1,cex.axis=1.5)

```



```

> fitq<- qda(group ~ ebitass + rotc,prior=c(9,1)/10,CV=T)
> fitq$class

```

```
[1] 1 1 1 1 1 1 1 1 1 1 1 1 2 2 2 2 2 2 1 2 2 2 2
Levels: 1 2
```

```
> fitq$posterior[1:10,]
```

```
      1      2
1 0.99996 3.7200e-05
2 1.00000 1.3842e-11
3 1.00000 3.0515e-13
4 1.00000 1.2232e-25
5 1.00000 1.8239e-10
6 1.00000 1.1134e-20
7 0.99678 3.2184e-03
8 1.00000 3.0345e-22
9 0.65731 3.4269e-01
10 1.00000 7.4754e-07
```

```
> table(group,fitq$class )
```

```
group  1  2
      1 12  0
      2  1 11
```

```
> #####
> #Support vector machines with ebitass data
> #####
>
> library(e1071)
> library(rpart)
> ddiscr <- read.table("C:/R/rmmva/data STEPDISCR SHARMA.txt", header=T, quote="\")
> attach(ddiscr)
> svm.model <- svm(as.factor(group) ~ rotc+ ebitass, data = ddiscr, cost = 100, gamma = 1)
> svm.pred <- predict(svm.model, ddiscr)
> table(svm.pred,ddiscr[,1])
```

```
svm.pred  1  2
          1 12  0
          2  0 12
```

```
> #####
> #Support vector machines with with rating data
> #####
>
> library(e1071)
> library(rpart)
> ddiscrm <- read.table("C:/R/rmmva/data discrim in R.txt", header=T, quote="\")
> attach(ddiscrm)
> #cost = the regularization parameter
> #gamma = parameter in kernel
> svm.model <- svm(as.factor(RATING) ~ LOPMAR+ LFIxCHAR +LGEARRAT+ LTDCAP+ LLEVER
+          + LCASHRAT + LACIDRAT +LCURRAT +LRECTURN +LASSLTD,
+          data = ddiscrm, cost = 100, gamma = 1)
> svm.pred <- predict(svm.model, ddiscrm)
> svm.model
```

```
Call:
svm(formula = as.factor(RATING) ~ LOPMAR + LFIxCHAR + LGEARRAT + LTDCAP +
    LLEVER + LCASHRAT + LACIDRAT + LCURRAT + LRECTURN + LASSLTD, data = ddiscrm,
    cost = 100, gamma = 1)
```

```
Parameters:
  SVM-Type:  C-classification
  SVM-Kernel: radial
    cost: 100
   gamma: 1
```

```
Number of Support Vectors: 95
```

```
> svm.pred
```

```

 1  2  3  4  5  6  7  8  9 10 11 12 13 14 15 16 17 18 19 20 21 22
AAA AAA AAA AAA AAA AAA AAA AAA AAA AAA AAx AAx AAx AAx AAx AAx AAx AAx AAx AAx AAx
23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40 41 42 43 44
AAx AAx AAx AAx Axx Axx Axx Axx Axx Axx Axx Axx Axx Axx Axx Axx Axx Axx BAA BAA BAA BAA
45 46 47 48 49 50 51 52 53 54 55 56 57 58 59 60 61 62 63 64 65 66
BAA BAA BAA BAA BAA BAA BAA BAA BAA BAA BAA BAx BAx BAx BAx BAx BAx BAx BAx BAx BAx BAx
67 68 69 70 71 72 73 74 75 76 77 78 79 80 81 82 83 84 85 86 87 88
BAx BAx BAx BAx Bxx Bxx Bxx Bxx Bxx Bxx Bxx Bxx Bxx Bxx Bxx Bxx Bxx Bxx C C C C C
89 90 91 92 93 94 95
C C C C C C C
Levels: AAA AAx Axx BAA BAx Bxx C
```

```
> table(svm.pred,ddiscrm[,1])
```

```
svm.pred AAA AAx Axx BAA BAx Bxx C
AAA 11 0 0 0 0 0 0
AAx 0 15 0 0 0 0 0
Axx 0 0 14 0 0 0 0
BAA 0 0 0 15 0 0 0
BAx 0 0 0 0 15 0 0
Bxx 0 0 0 0 0 13 0
C 0 0 0 0 0 0 12
```

```
> #####
> #classification trees
> #####
> par(mar=c(0.2, 0.2, 0.2, 0))
> par(oma=c(0, 3, 3, 0))
> workdataact <- read.table("C:/R/rmmva/workdata.txt", header=T, quote="\")
> attach(workdataact)
> #classification trees
> library(rpart)
> tree <- rpart(thisyr ~ child1 +child2 +black +hubinc+ educ+ age,parms=list(split="gini"))
> tree
```

```
n= 100
```

```
node), split, n, deviance, yval
```

* denotes terminal node

```
1) root 100 20.1600 0.72000
 2) age< 34.5 47 11.4890 0.57447
   4) hubinc>=10.885 11 2.1818 0.27273 *
   5) hubinc< 10.885 36 8.0000 0.66667
      10) hubinc< 7.9645 26 6.1538 0.61538
         20) age>=30.5 9 2.2222 0.44444 *
         21) age< 30.5 17 3.5294 0.70588 *
      11) hubinc>=7.9645 10 1.6000 0.80000 *
 3) age>=34.5 53 6.7925 0.84906
   6) hubinc>=3.895 43 6.5116 0.81395
      12) hubinc< 6.486 9 2.2222 0.55556 *
      13) hubinc>=6.486 34 3.5294 0.88235
         26) hubinc>=11.296 15 2.9333 0.73333 *
         27) hubinc< 11.296 19 0.0000 1.00000 *
 7) hubinc< 3.895 10 0.0000 1.00000 *
```

```
> printcp(tree) #display the crossvalidation results
```

Regression tree:

```
rpart(formula = thisyr ~ child1 + child2 + black + hubinc + educ +
      age, parms = list(split = "gini"))
```

Variables actually used in tree construction:

```
[1] age    hubinc
```

Root node error: 20.2/100 = 0.202

n= 100

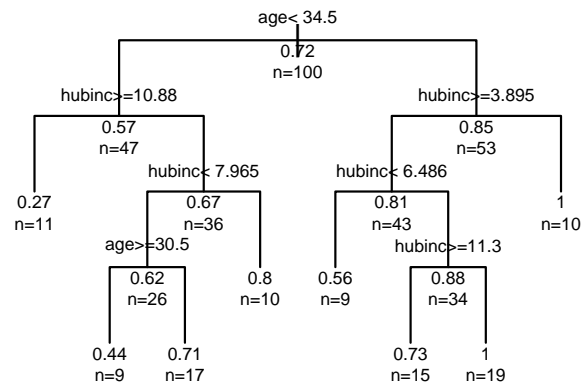
	CP	nsplit	rel error	xerror	xstd
1	0.0932	0	1.000	1.02	0.100
2	0.0649	1	0.907	1.07	0.119
3	0.0258	2	0.842	1.09	0.123
4	0.0161	5	0.761	1.21	0.136
5	0.0100	7	0.729	1.22	0.137

```
> # summary(tree) #display more details about the tree construction
```

```
> plot(tree,uniform = TRUE, margin = 0.20,main ="working mom classification tree" )
```

```
> text(tree, use.n = TRUE, all=TRUE, cex=0.5)
```


working mom classification tree



```
> smallertree <- prune(tree,cp=0.041)
> smallertree
```

n= 100

```
node), split, n, deviance, yval
      * denotes terminal node
```

```
1) root 100 20.1600 0.72000
  2) age< 34.5 47 11.4890 0.57447
    4) hubinc>=10.885 11 2.1818 0.27273 *
    5) hubinc< 10.885 36 8.0000 0.66667 *
  3) age>=34.5 53 6.7925 0.84906 *
```