Code application PLS-DA

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```
knitr::opts_chunk$set(echo = TRUE)

rm(list = ls())
```

CHARGEMENT PACKAGE

```
library(plsdepot)
library(ggplot2)
library(ggrepel)
```

CHARGEMENT FONCTION CIRCLE FUN

```
setwd("~/Presentations/PLS-DA") ## Ouverture session avec chemin fonction
source("CircleFun.r")
```

CHARGEMENT FICHIER DE DONNEES

```
setwd("~/Presentations/PLS-DA") ## Ouverture session avec chemin jeu de donnees
load(file = "data_PLSDA.RData")
## Jeu de donnees :
## 3 premieres colonnes = description observations
## colonne 4 a 44 = 41 variables explicatives
## colonne 45 = variable a expliquer (lame_score)
```

MISE EN FORME DES VARIABLES

```
## CREATION DE LA MATRICE DE VARIABLES X
Xvar <- data_select[,4:(ncol(data_select)-1)]
## suppression variables qui decrivent les observations et variable a predire

## CREATION DE LA MATRICE DES VARIABLES Y DICHOTOMISEE
no_lame <- ifelse(data_select$lame_score=="no_lame",1,0)
## si observation non boiteuse, alors 1 sinon 0
lame = ifelse(data_select$lame_score=="lame",1,0)</pre>
```

```
## si observation boiteuse, alors 1 sinon 0

Yvar <- data.frame(no_lame = no_lame, lame = lame)

rm(no_lame)
rm(lame)</pre>
```

APPLICATION DE LA PLS-DA (PLS-2 SUR VARIABLE Y DI-CHOTOMISEE)

```
res.plsreg2 <- plsreg2(Xvar, Yvar, comps = 10, crosval = TRUE)
## remarque : par defaut, centrage et reduction variables par fonction plsreg2
```

CHOIX DU NOMBRE DE COMPOSANTES A RETENIR

```
## Selection a partir evolution erreur classification
## obtenue par cross validation selon nombre de composantes
## ou selection variables avec Q2 :
## critere de Tenenhaus : composante h retenue si Q2h > 0.0975
## (ou Q2h > 0.05 selon reference)
res.plsreg2$Q2 ## ici seulement premiere composante significative
```

```
## t1 Q2.no_lame Q2.lame Q2

## t1 0.11424141 0.11424141 0.11424141

## t2 0.02533965 0.02533965 0.02533965

## t3 -0.01819481 -0.01819481 -0.01819481

## t4 -0.02544873 -0.02544873 -0.02544873

## t5 -0.05124956 -0.05124956 -0.05124956

## t6 -0.04139387 -0.04139387 -0.04139387

## t7 -0.03947761 -0.03947761 -0.03947761

## t8 -0.04640724 -0.04640724 -0.04640724

## t9 -0.04751747 -0.04751747 -0.0495937
```

res.plsreg2\$Q2cum ## mais amelioration Q2cum deuxieme composante :

```
##
     Q2cum.no_lame Q2cum.lame
                                Q2cum
## t1
        0.11424141 0.11424141 0.11424141
## t2
        ## t3
        0.12097839 0.12097839 0.12097839
        0.09860841 0.09860841 0.09860841
## t4
## t5
      0.05241248 0.05241248 0.05241248
## t6
      ## t7
       -0.02576881 -0.02576881 -0.02576881
## t8
      -0.07337190 -0.07337190 -0.07337190
## t9
      -0.12437583 -0.12437583 -0.12437583
## t10
      -0.18054894 -0.18054894 -0.18102653
```

```
## 1ere et 2eme composantes retenues --> projection plan (t1, t2) OK, avec ## information essentiellement restituee sur t1
```

PERFORMANCE DU MODELE DE LA PLS-DA

```
Q2cum <- round(res.plsreg2$Q2cum[2]*100) ## Q2 cum deux premieres compo
R2xcum <- round(res.plsreg2$expvar[2,2]*100) ## R2xcum deux premieres compo
R2ycum <- round(res.plsreg2$expvar[2,4]*100) ## R2ycum cum deux premieres compo
## Remarque : R2 et Q2 assez faibles
## --> modele developpe non predictif, mais 1ere composante
## significative --> une partie de l'information de Y (17% >> 0%)
## effectivement expliquee par X
```

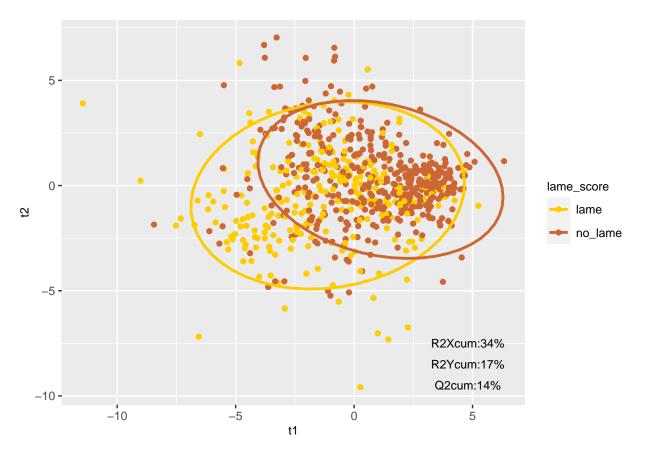
IDENTIFICATION DES VARIABLES LES PLUS DISCRIMINANTES AVEC LES VIP

```
## Variables retenues --> VIP > 0.8 sur les deux premieres composantes
var_imp <-res.plsreg2$VIP[which(res.plsreg2$VIP[,2]>=0.8),2]
var_imp_names <-rownames(res.plsreg2$VIP[which(res.plsreg2$VIP[,2]>=0.8),])

var_imp_sort <- sort(var_imp,decreasing = TRUE) ## Variables triees par ordre
## d'importance decroissant a partir des valeurs des VIP
## sur les deux premieres composantes</pre>
```

VISUALISATION DES INDIVIDUS ET VARIABLES SUR LES DEUX PREMIERES COMPOSANTES PLS

```
## Plot des individus
df_pda_ind <- data.frame(score_t1 = res.plsreg2$x.scores[,1],</pre>
                         score_t2 = res.plsreg2$x.scores[,2],
                         lame score = data select$lame score)
p <- ggplot(data = df_pda_ind, aes(x = score_t1, y = score_t2, col = lame_score))+
geom_point() + annotate(geom="text", x=4.8, y=-7.5,
label=paste("R2Xcum:", R2xcum, "%", sep = ""), color="black", size = 3) +
annotate(geom="text", x=4.8, y=-8.5,
label=paste("R2Ycum:", R2ycum, "%", sep = ""), color="black", size = 3) +
annotate(geom="text", x=4.8, y=-9.5,
label=paste("Q2cum:", Q2cum, "%", sep = ""), color="black", size = 3) +
scale_color_manual(values=c("#FFCC00", "#CC6633"))+ stat_ellipse(size = 1) +
theme(axis.title.x = element_text(size = 9),
axis.title.y = element_text(size = 9),legend.title = element_text(size = 9),
legend.text = element text(size = 9)) +
xlab("t1") + ylab("t2")
```



```
## Plot du cercle des correlations avec les variables selectionnees
df_pda_var <-
  res.plsreg2$cor.xt[which(rownames(res.plsreg2$cor.xt)%in%var_imp_names),1:2]
df_pda_var<- data.frame(rbind(df_pda_var, res.plsreg2$cor.yt[,1:2]))</pre>
df_pda_var$type <- c(rep("variable", (nrow(df_pda_var)-2)), rep("lame_score",2))</pre>
df_pda_var$type <- as.factor(df_pda_var$type)</pre>
df_pda_var=cbind(x1=rep(0,times=dim(df_pda_var)[1]),
                 x2=rep(0,times=dim(df_pda_var)[1]),
                 df_pda_var)
colnames(df_pda_var)=c("x1","y1","xend","yend","type")
dat <- circleFun(npoints = 1000)</pre>
my_color <- ifelse(df_pda_var$type == "variable","#009999","#FF9966")</pre>
g=ggplot(dat,aes(x,y)) + geom_path()+
scale_x_continuous(breaks=seq(-1, 1, by=1))+
scale_y_continuous(breaks=seq(-1, 1, by=1))
g=g + theme_bw()+theme(panel.grid.minor = element_blank())
g=g + geom_segment(aes(x = x1, y = y1, xend = xend, yend = yend),color="#009999",
data = df_pda_var,
arrow = arrow(length = unit(0.01, "npc"))) + xlab("t1") + ylab("t2") +
geom_label_repel(data = df_pda_var,
aes(xend, yend, label = rownames(df_pda_var)), size = 2,
inherit.aes=FALSE,color = my_color,fill ="white",arrow=arrow(length(unit(0,'inches'))),
box.padding = 0)
g
```

