

Homework2: Dim-Reduction

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1. PAM only

PAM is included in cluster package.

```
# Require package
library(cluster)
library(mclust)

## Package 'mclust' version 5.4.5
## Type 'citation("mclust")' for citing this R package in publications.

# input data
data.snp <- read.table("~/Codes/GWAS/HW2/c1_snps_rec1.txt", header = TRUE)

# Creat a table between the true origins and the clustering result
data.type <- rep("Asian", 697)
data.type[data.snp$racess %in% c("CEPH - 1",
                                "CEPH - 2",
                                "Tuscan",
                                "Tuscan - Additional")] <- "European"
data.type[data.snp$racess %in% c("Luhya",
                                "Luhya - Additional",
                                "Yoruba - 1",
                                "Yoruba - 2",
                                "Yoruba - Additional")] <- "African"

# Clustering by PAM with k = 3
fit.pam <- pam(data.snp[, -c(1,2)], k = 3, metric = "euclidean")
cluster.pam <- fit.pam$clustering

# Classification error rate
error.PAM <- classError(cluster.pam, data.type)$errorRate
cat("The classification error rate of PAM on original data is: ", error.PAM, "\n")

## The classification error rate of PAM on original data is: 0.02582496

# table
table(cluster.pam, data.type)

##           data.type
## cluster.pam African Asian European
##           1      212      0        0
##           2        1     316        5
##           3         7       5     151
```

2. PAM after PCA

```
# Apply PCA
fit.pca <- prcomp(data.snp[, -c(1, 2)])
```

```

# Calculate proportion and cumulative proportion of variance explained by each PC
variance.table <- data.frame(Var = round(fit.pca$sdev^2),
                             Prop<- fit.pca$sdev^2/sum(fit.pca$sdev^2)*100,
                             Cum.Prop<- cumsum(fit.pca$sdev^2/sum(fit.pca$sdev^2)*100))
variance.table.round <- round(variance.table, digits = 3)
names(variance.table.round)[c(2,3)] <- c("Prop", "Cum.Prop")
head(variance.table.round, 10)

```

```

##      Var      Prop Cum.Prop
## 1      8 11.081   11.081
## 2      4  5.569   16.650
## 3      1  1.999   18.649
## 4      1  1.678   20.327
## 5      1  1.604   21.931
## 6      1  1.486   23.417
## 7      1  1.444   24.861
## 8      1  1.385   26.246
## 9      1  1.315   27.561
## 10     1  1.301   28.863

```

```

num.pc <- c(2, 5, 10)
for (num in num.pc) {
  pcs <- fit.pca$x[,1:num]
  # Clustering by PAM with k = 3
  fit.pam <- pam(pcs, k = 3, metric = "euclidean")
  cluster.pam <- fit.pam$clustering

  # Classification error rate
  error.PAM <- classError(cluster.pam, data.type)$errorRate
  cat("\nThe classification error rate of PAM on", num, "pcs is: \t", error.PAM, "\n")
  # table
  print(table(cluster.pam, data.type))
}

```

```

##
## The classification error rate of PAM on 2 pcs is:      0.01147776
##      data.type
## cluster.pam African Asian European
##           1      215      0      0
##           2       0    319      1
##           3       5     2    155
##
## The classification error rate of PAM on 5 pcs is:      0.01004304
##      data.type
## cluster.pam African Asian European
##           1      215      0      0
##           2       0    320      1
##           3       5     1    155
##
## The classification error rate of PAM on 10 pcs is:     0.01147776
##      data.type
## cluster.pam African Asian European
##           1      215      0      1
##           2       0    320      1

```

##	3	5	1	154
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