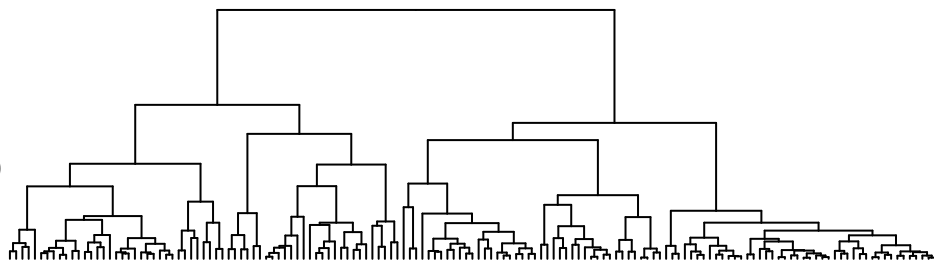
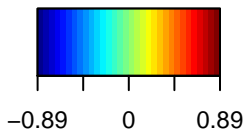


Color key



help("cim")

ALT.IU.L.

AST.IU.L.

BUN.mg.dL.

TBA.umol.L.

ALP.IU.L.

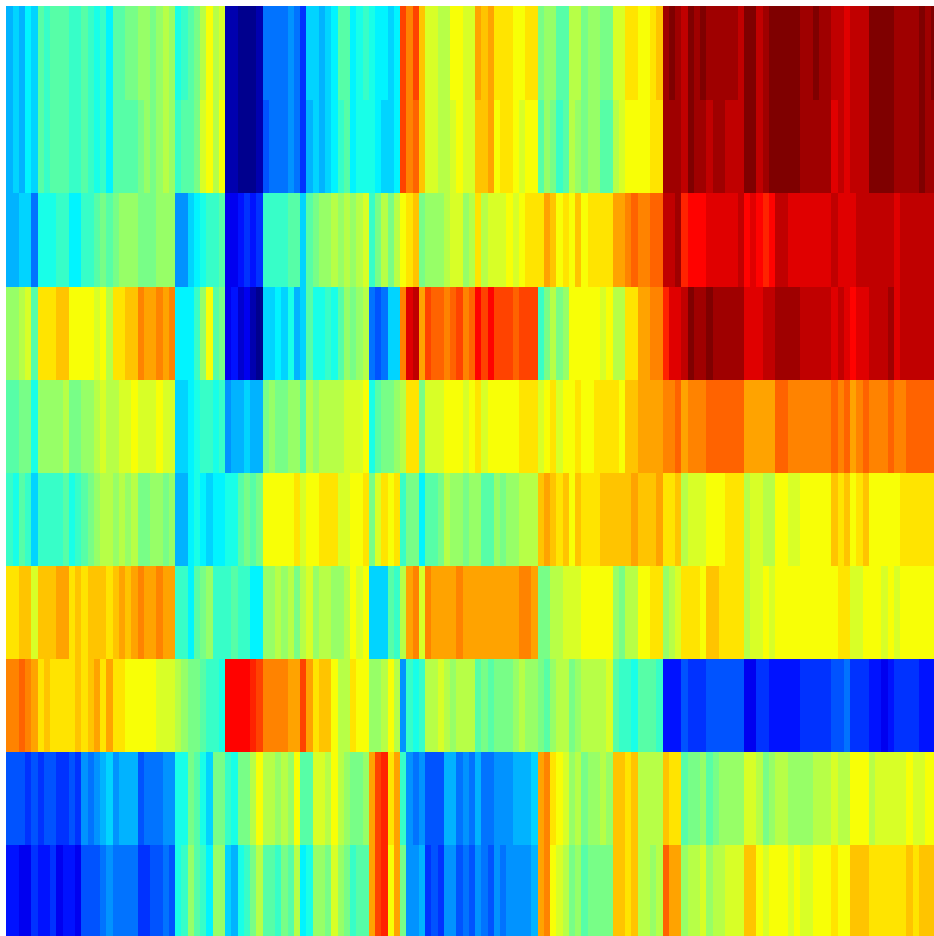
Creat.mg.dL.

SDH.IU.L.

Cholesterol.

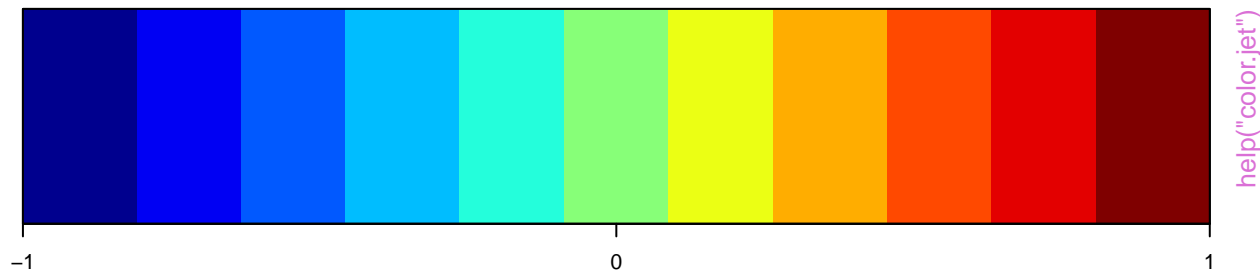
TP.g.dL.

ALB.g.dL.

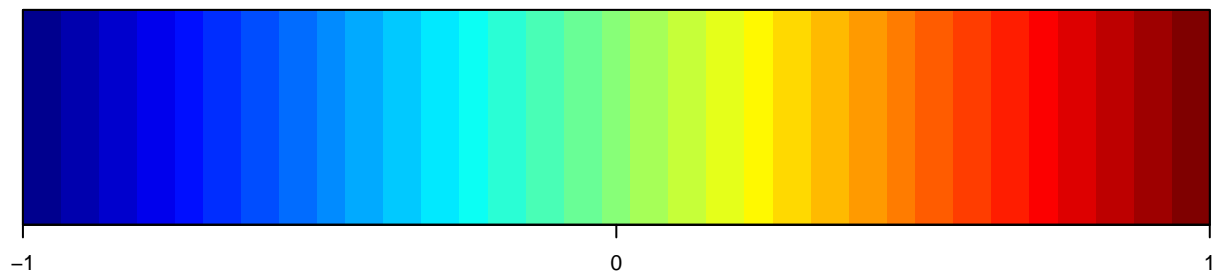


Gene expression data (rows) and sample labels (columns) are displayed below the heatmap.

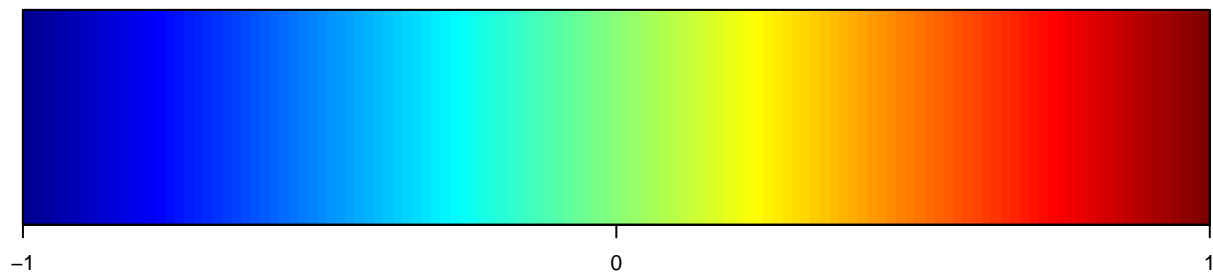
**n = 11**



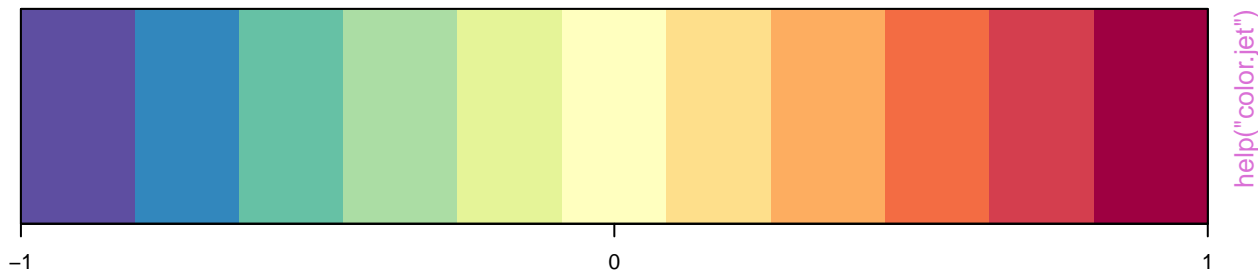
**n = 33**



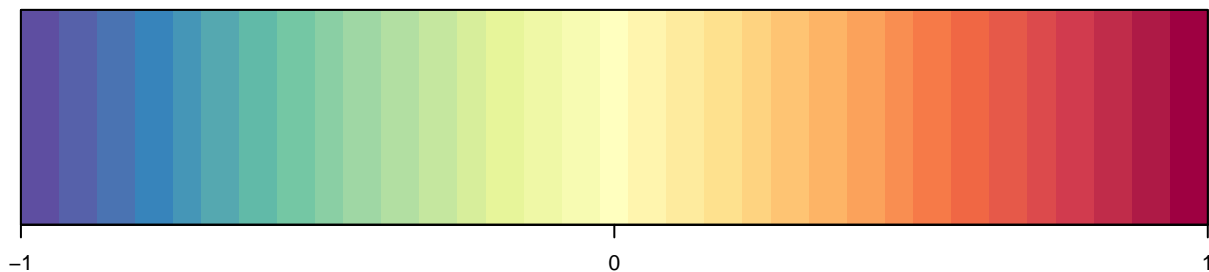
**n = 125**



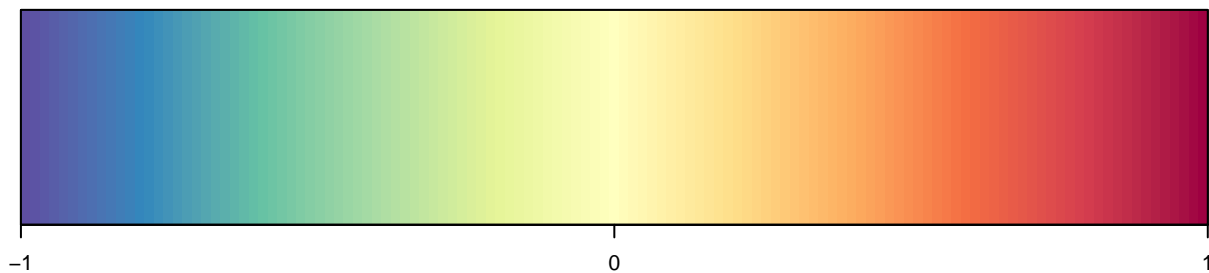
**n = 11**



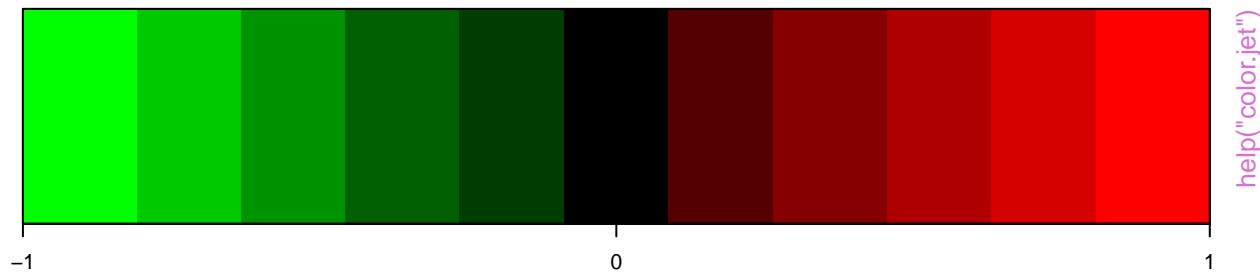
**n = 33**



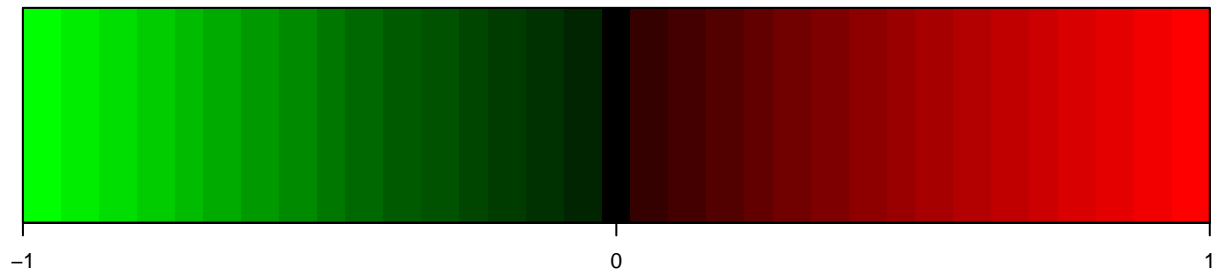
**n = 125**



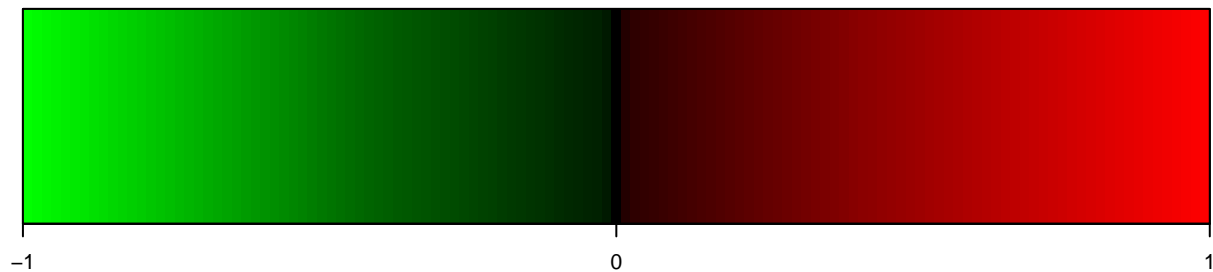
**n = 11**

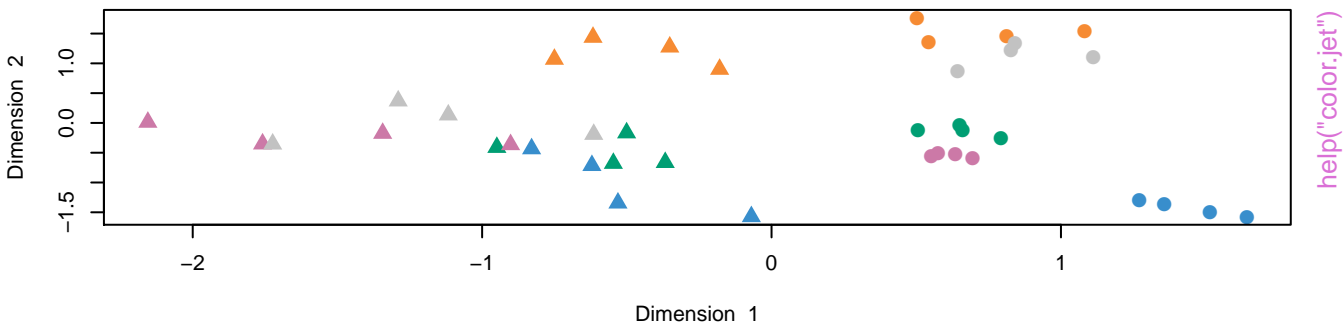


**n = 33**



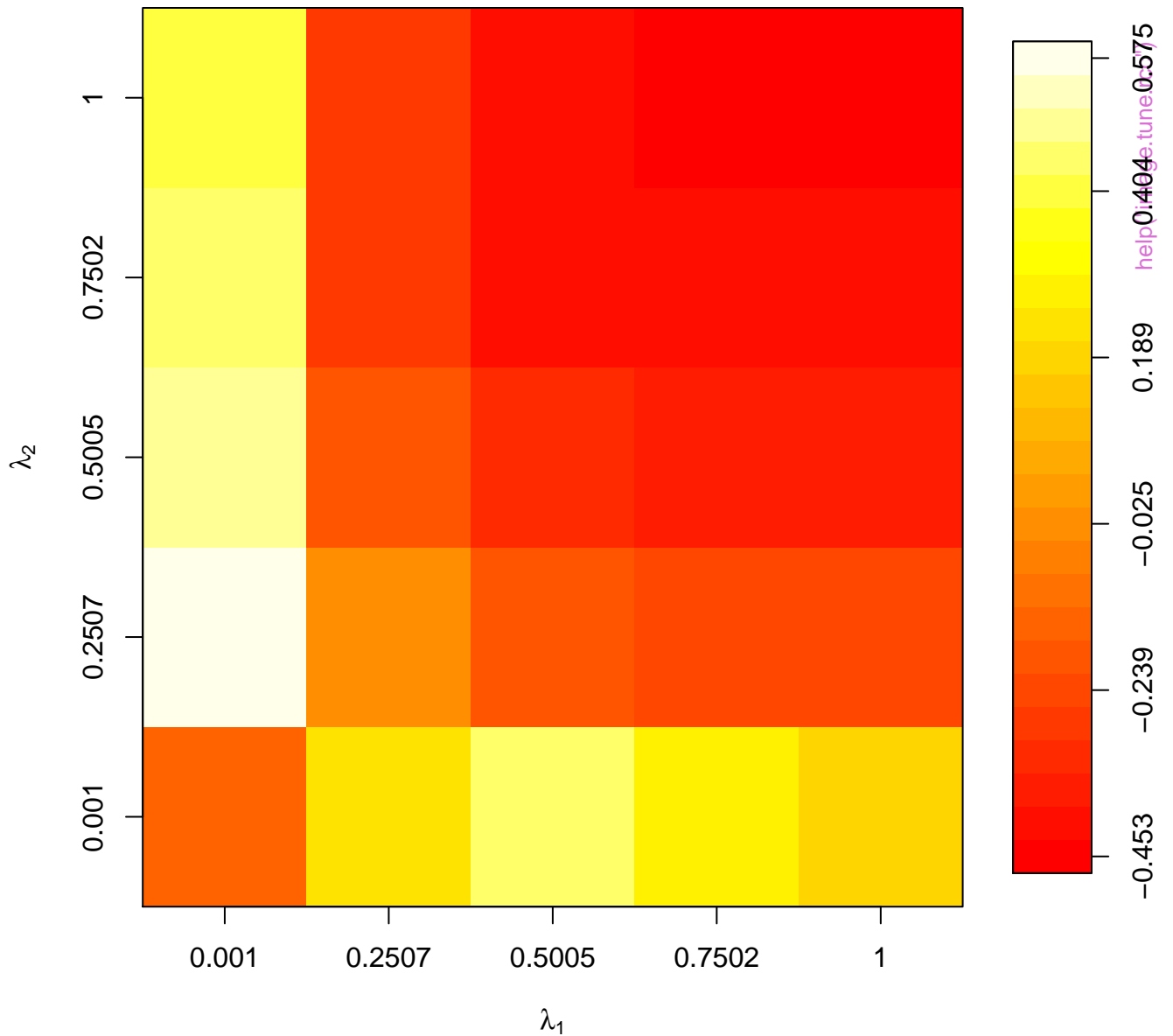
**n = 125**



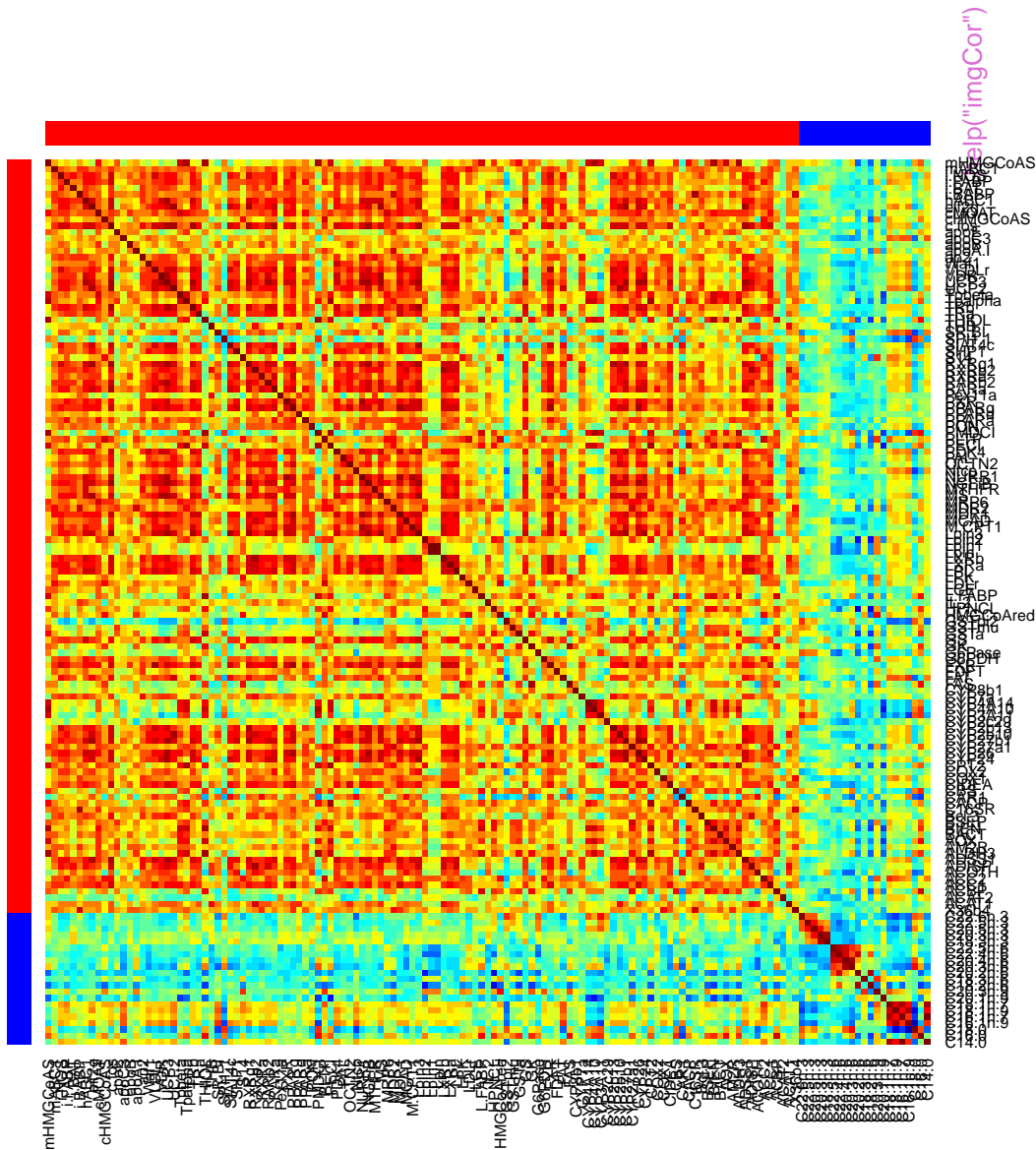
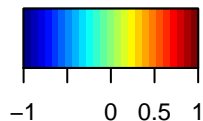


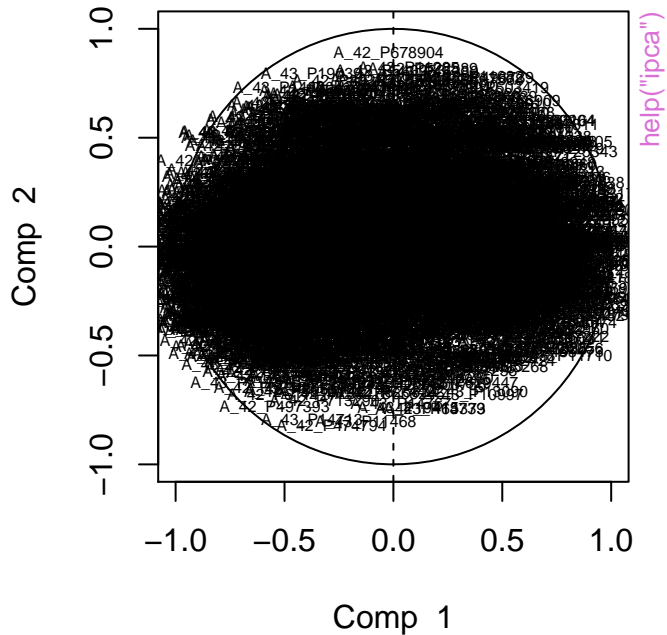
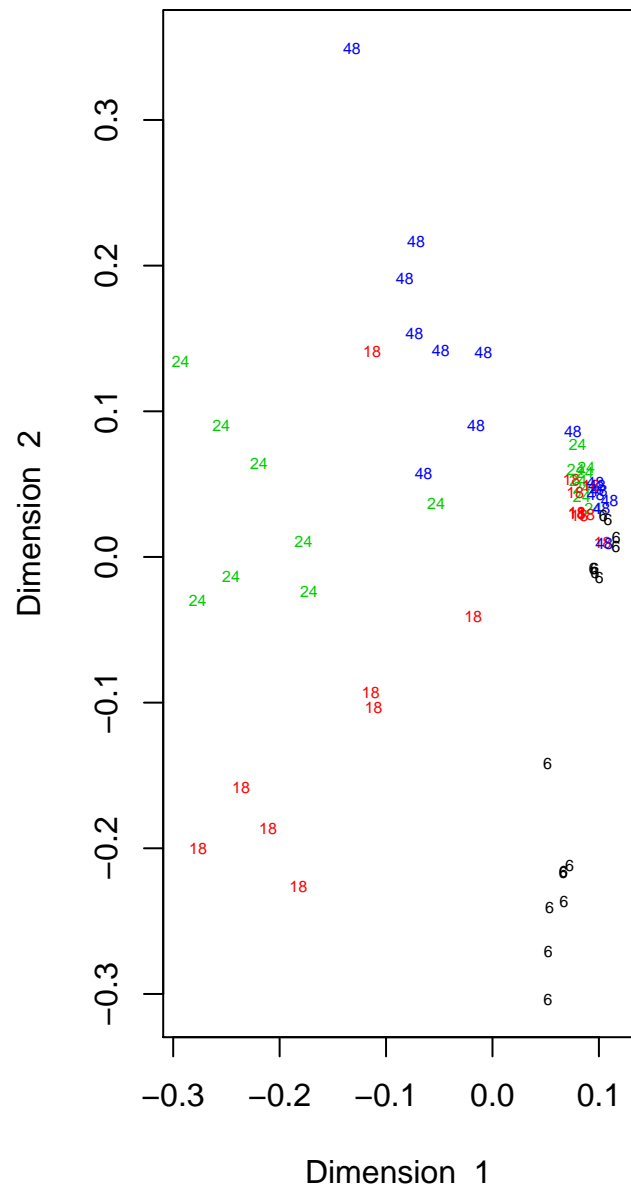


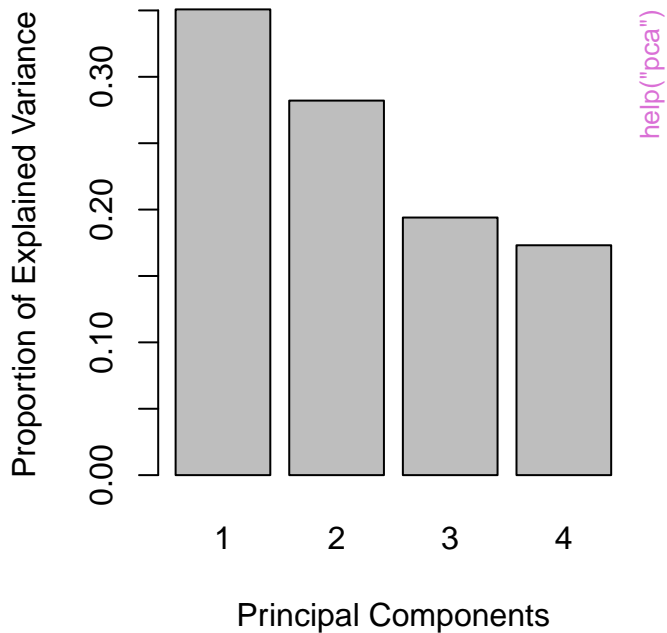
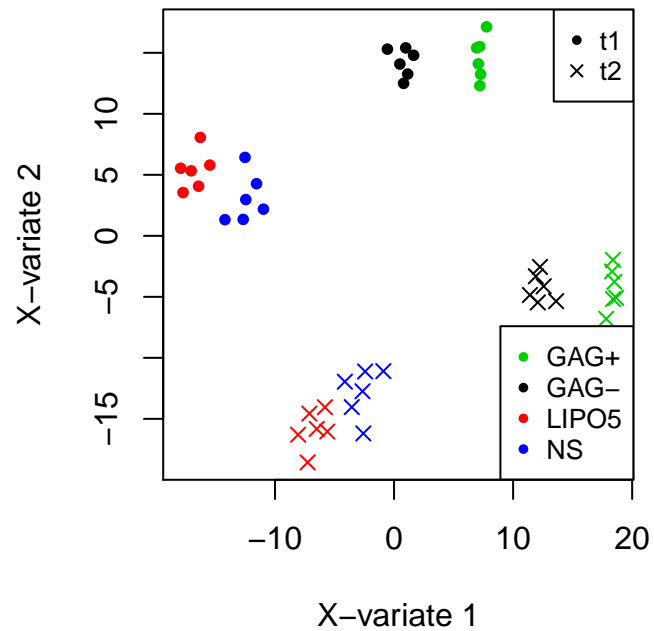
$$CV(\lambda_1, \lambda_2)$$

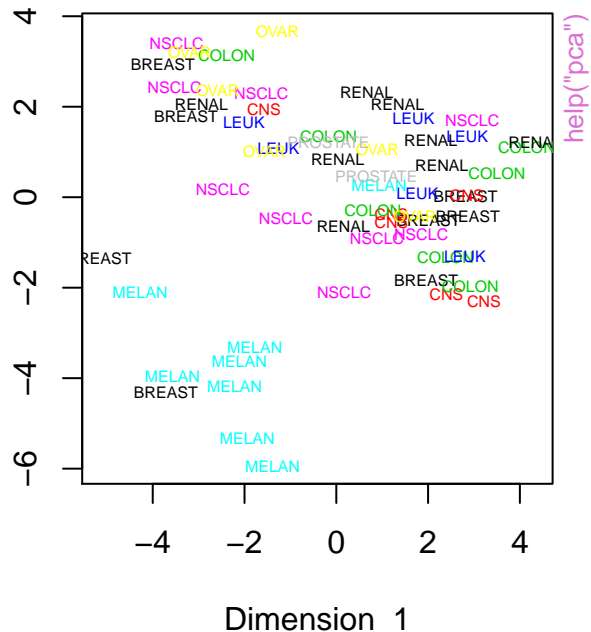
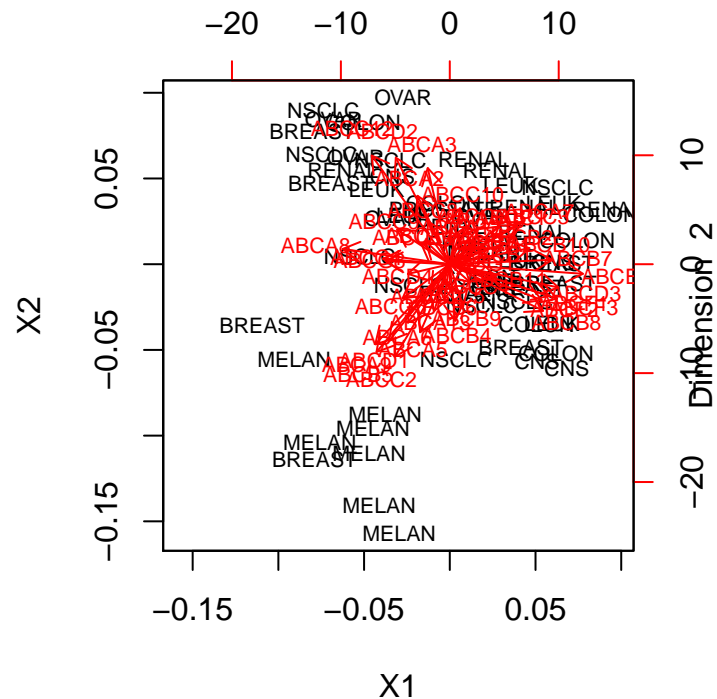


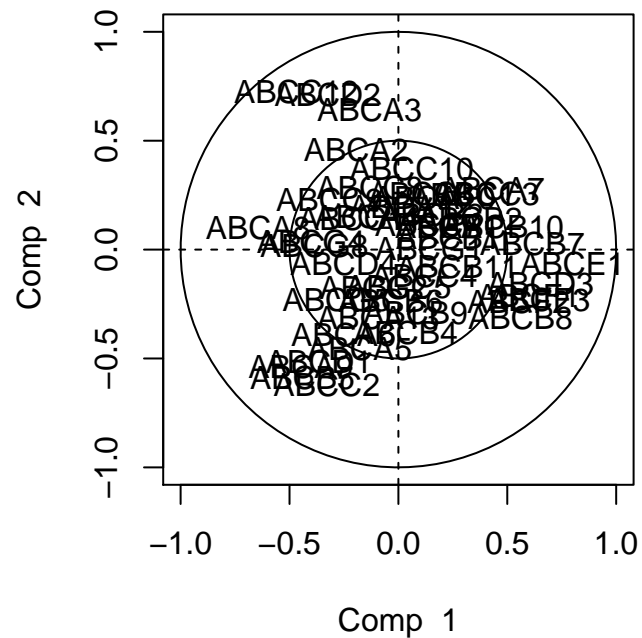
## [X,Y] correlation matrix

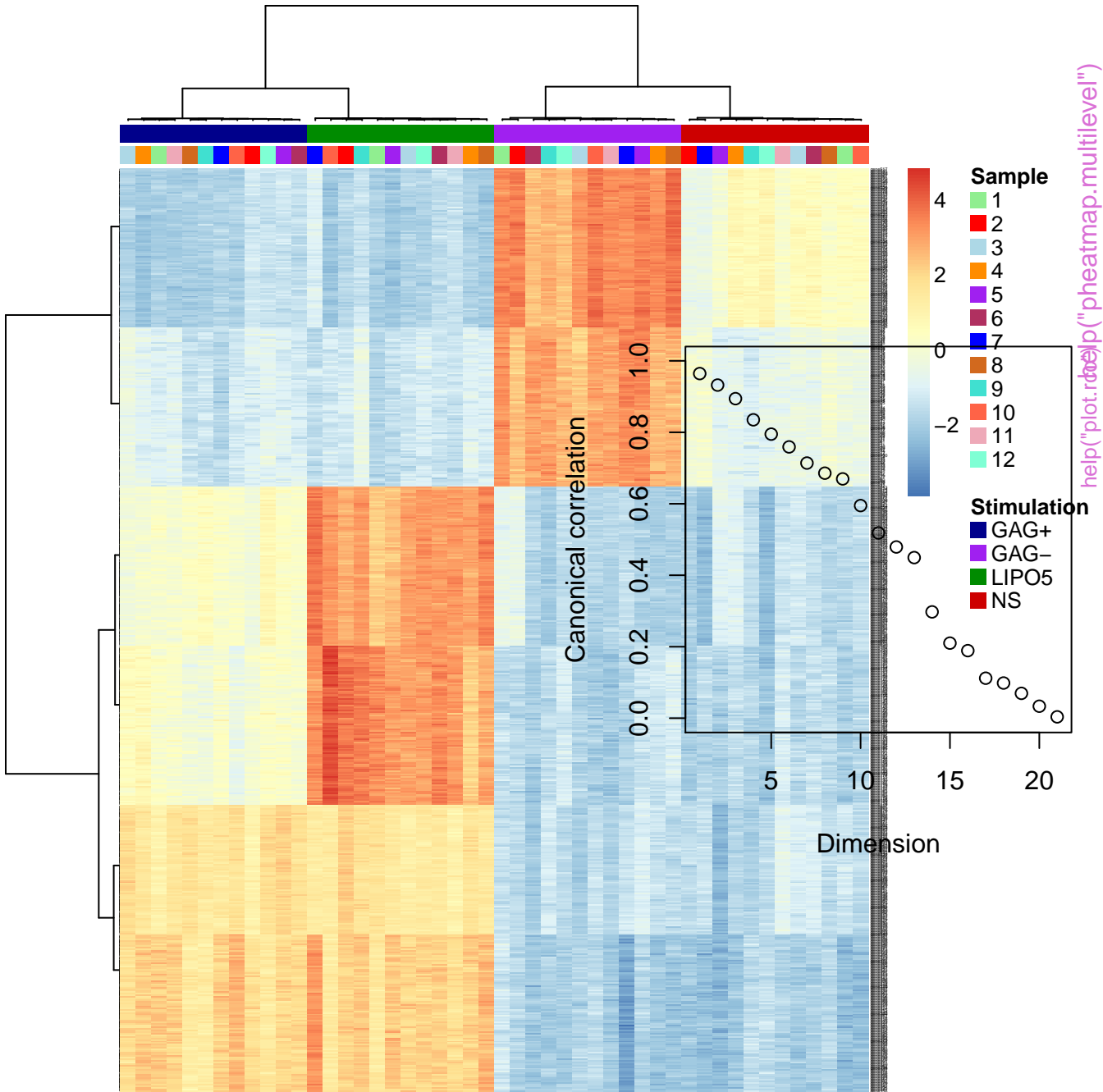


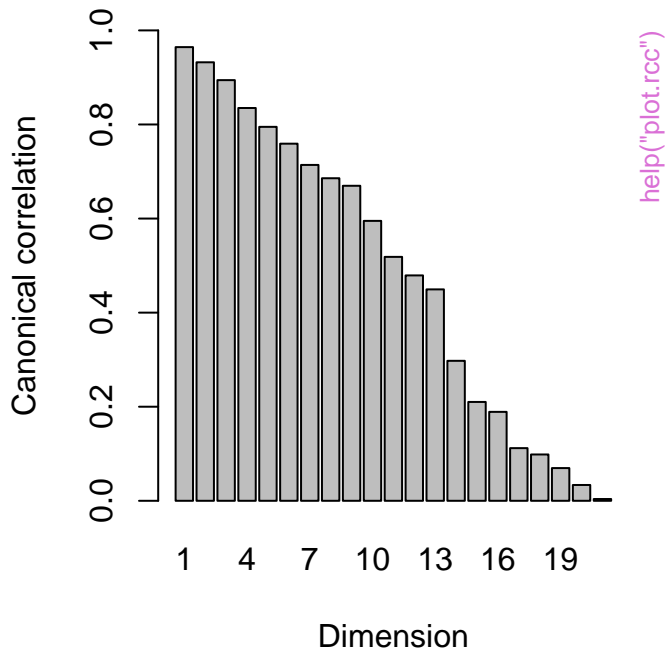
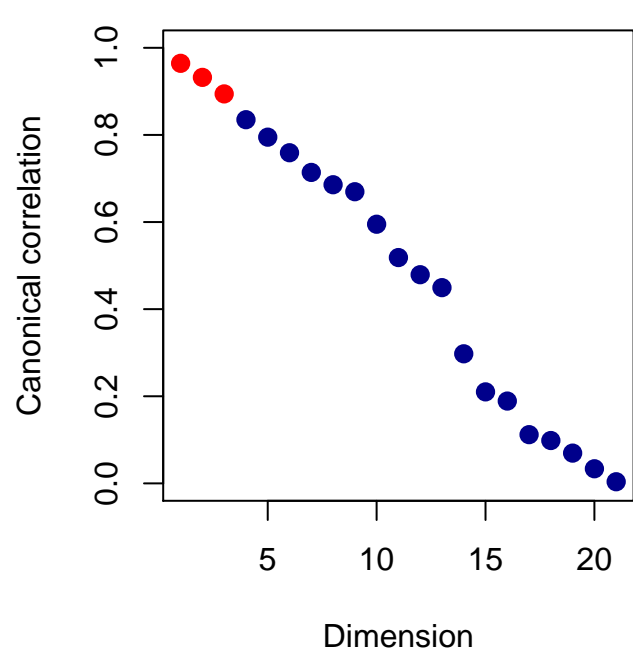




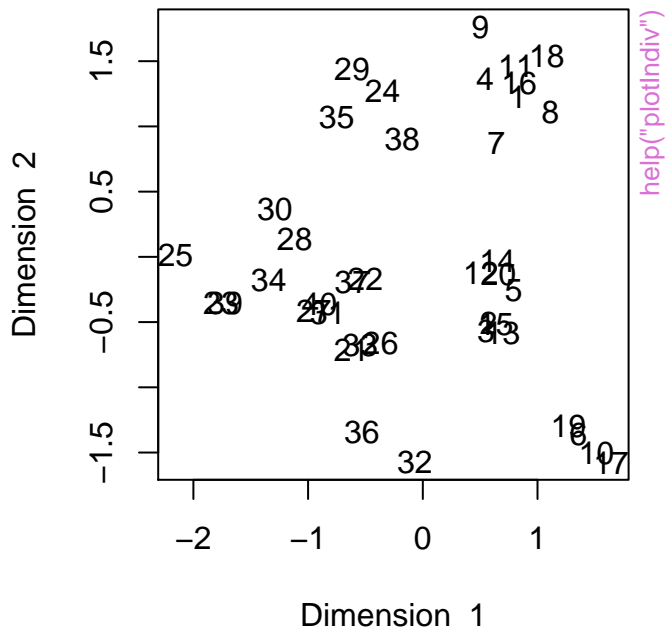
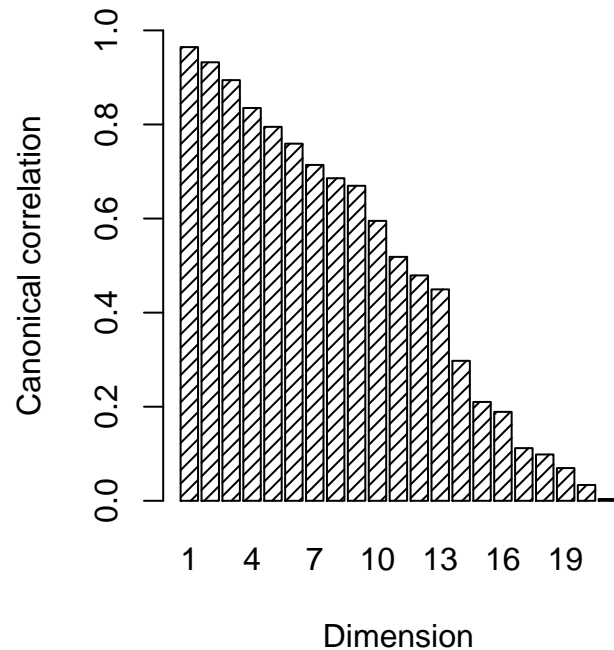


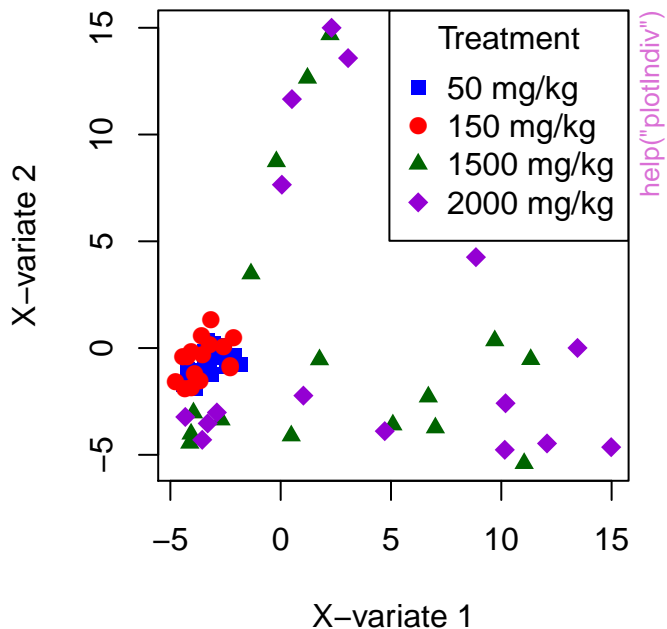
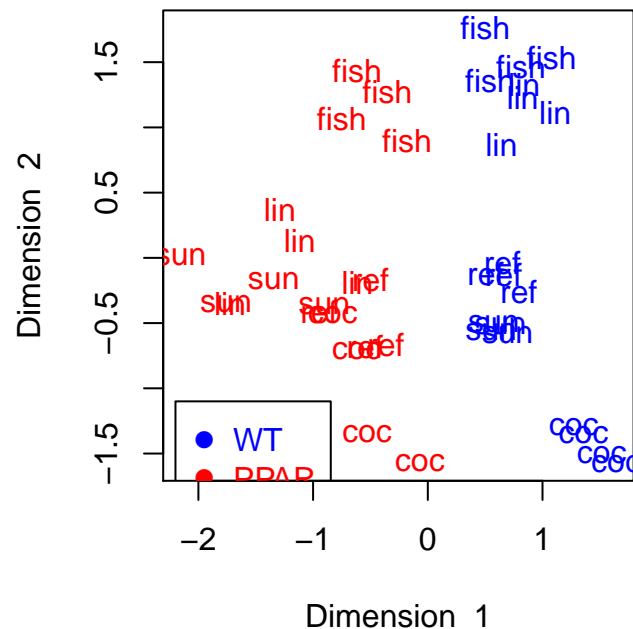


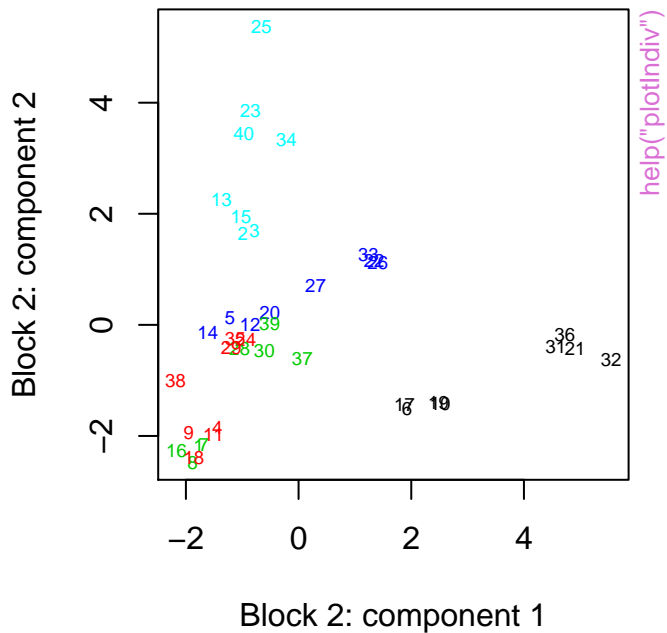
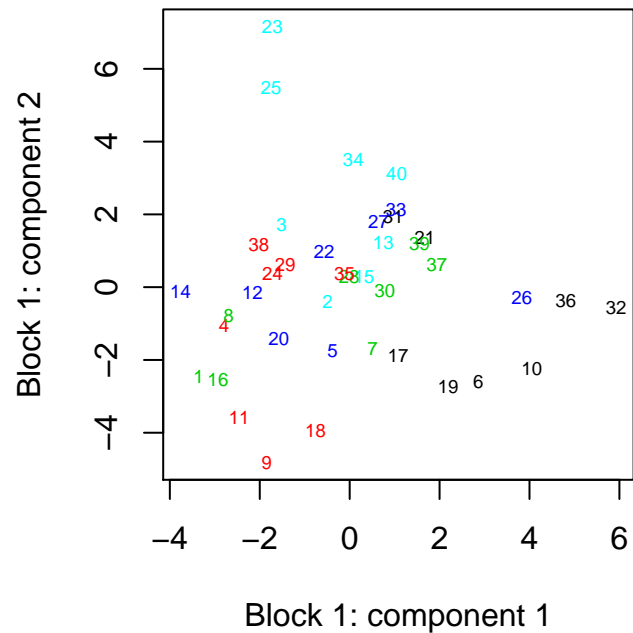


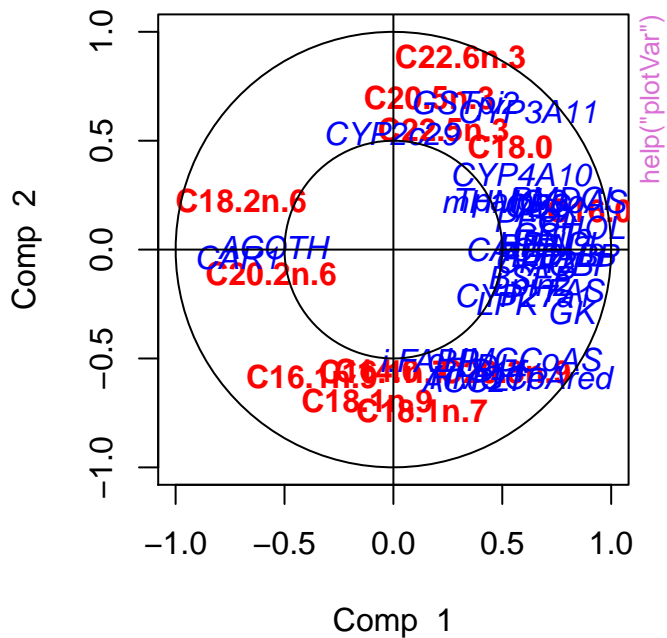
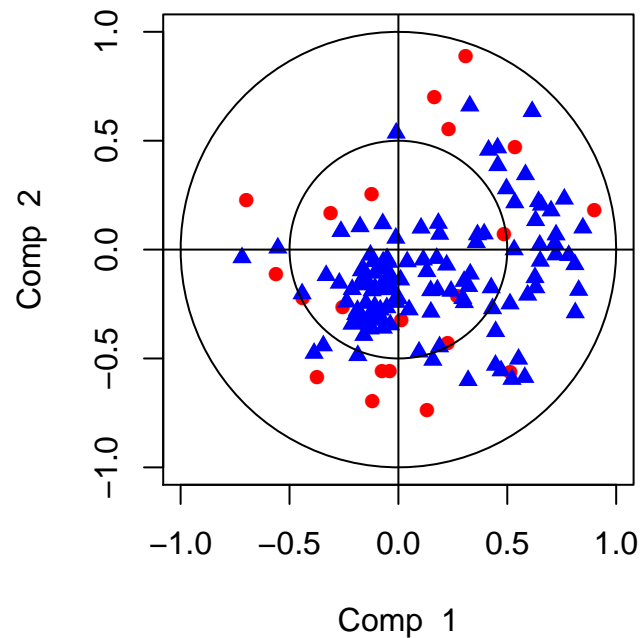


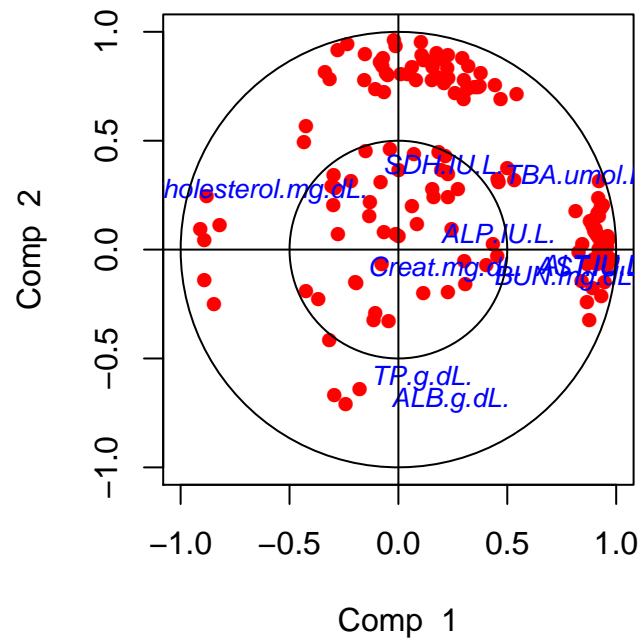




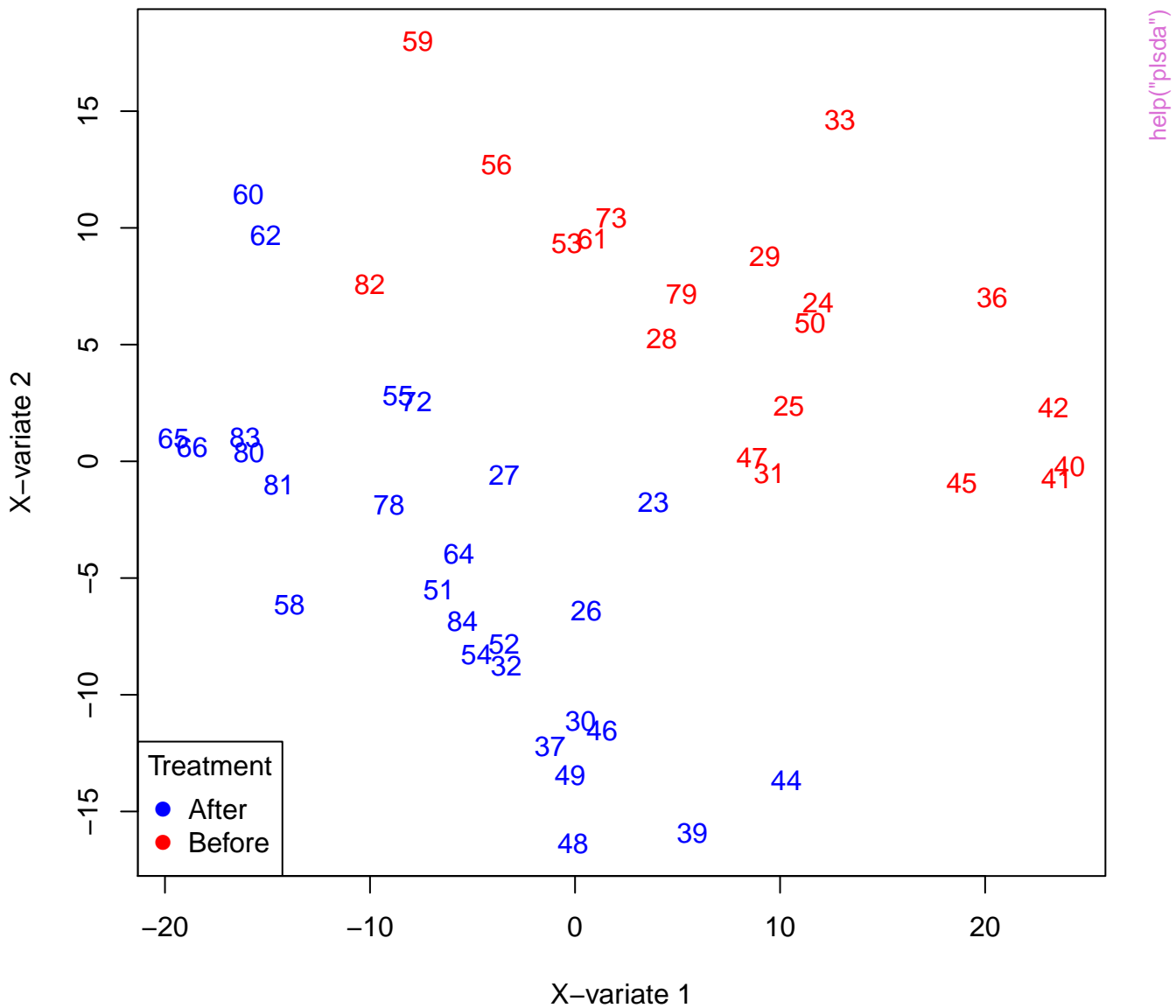


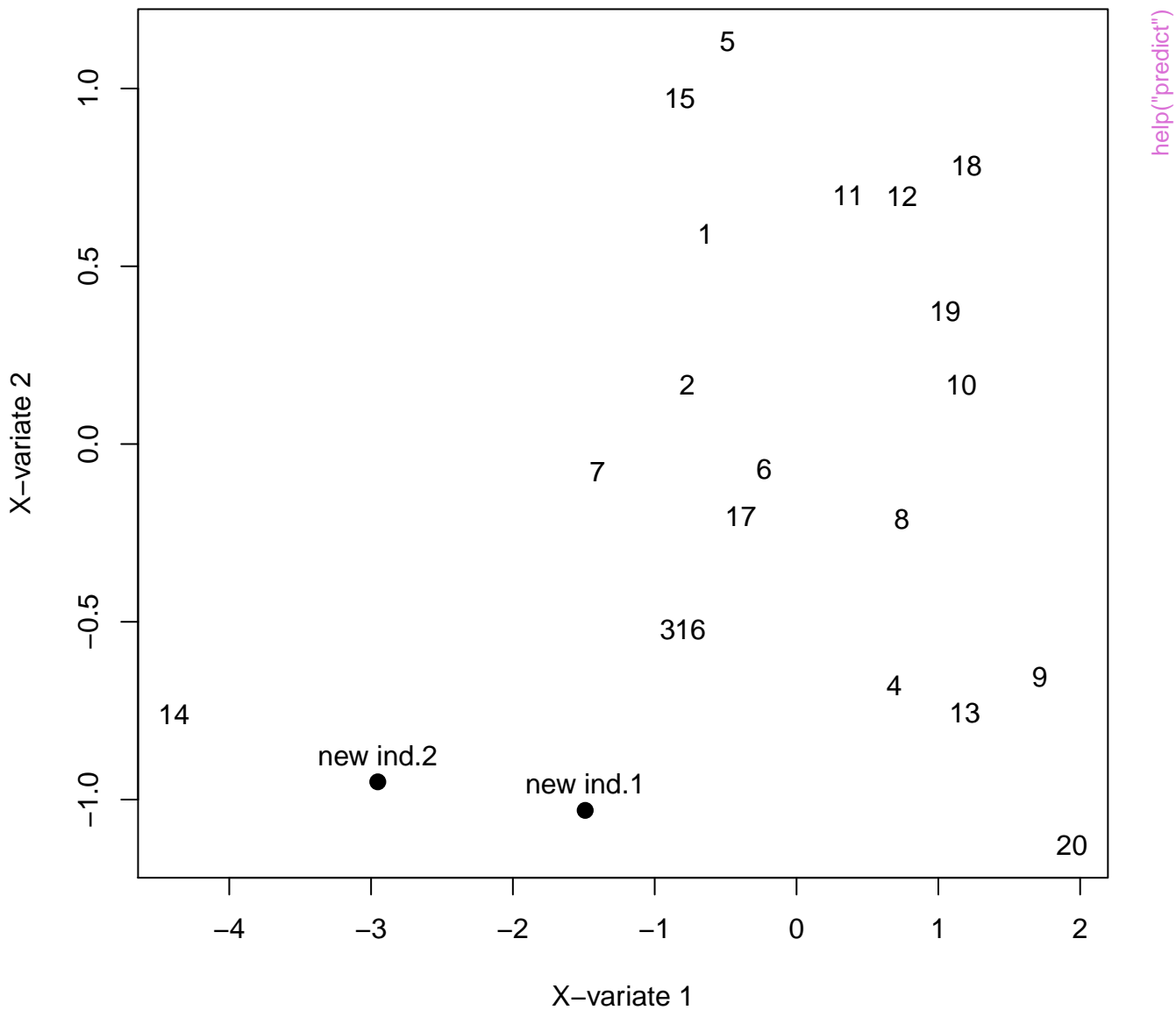




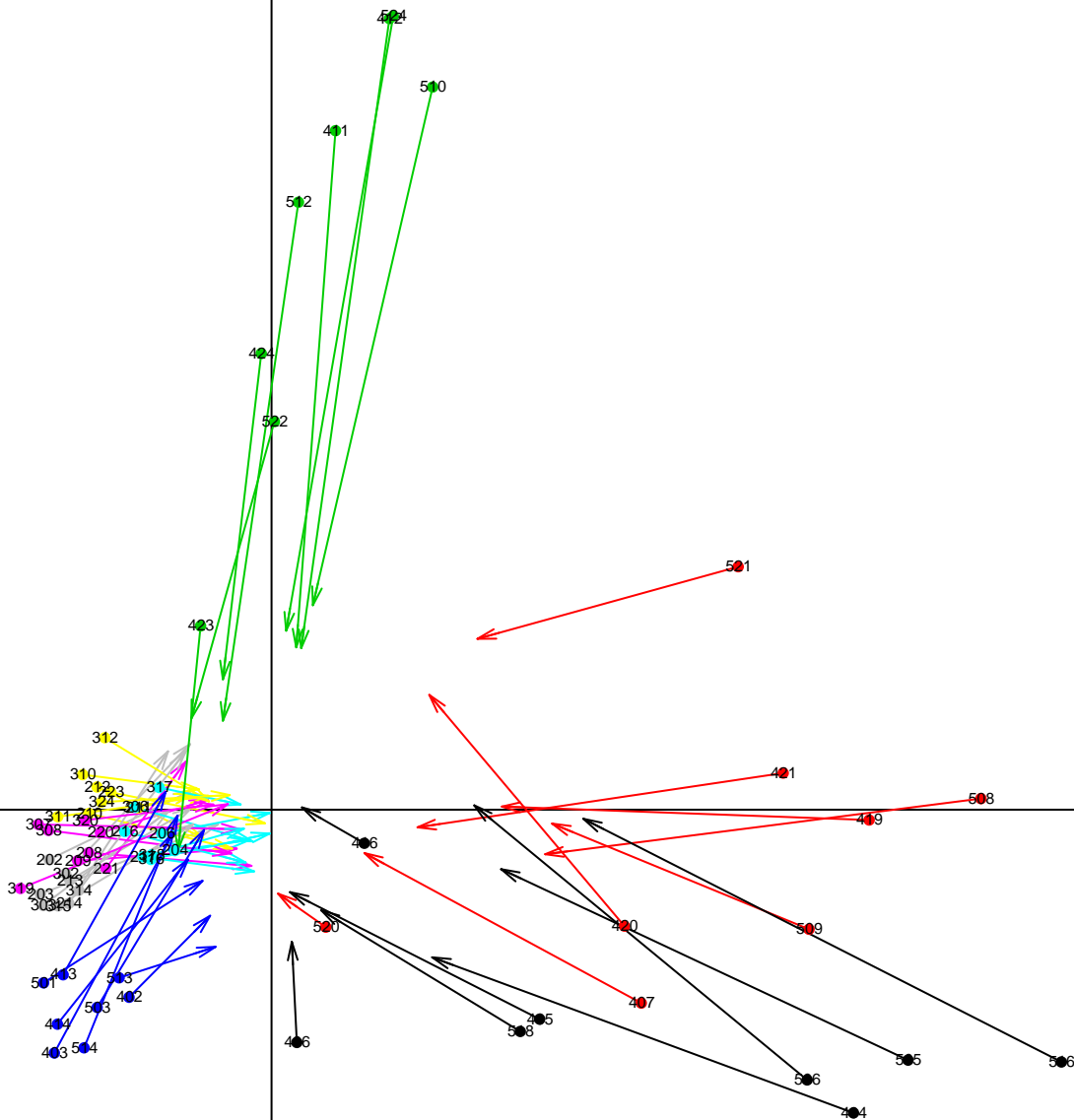




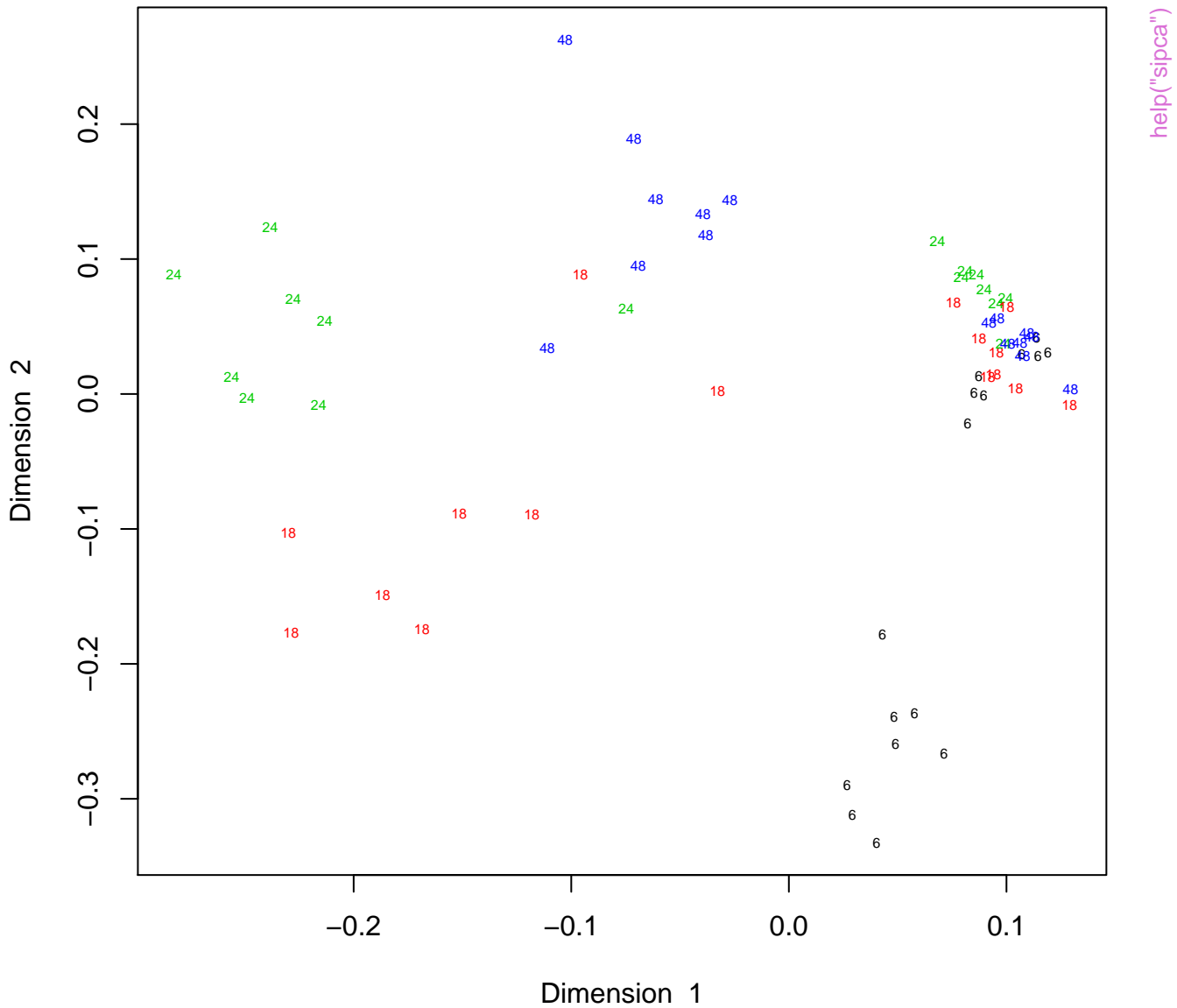




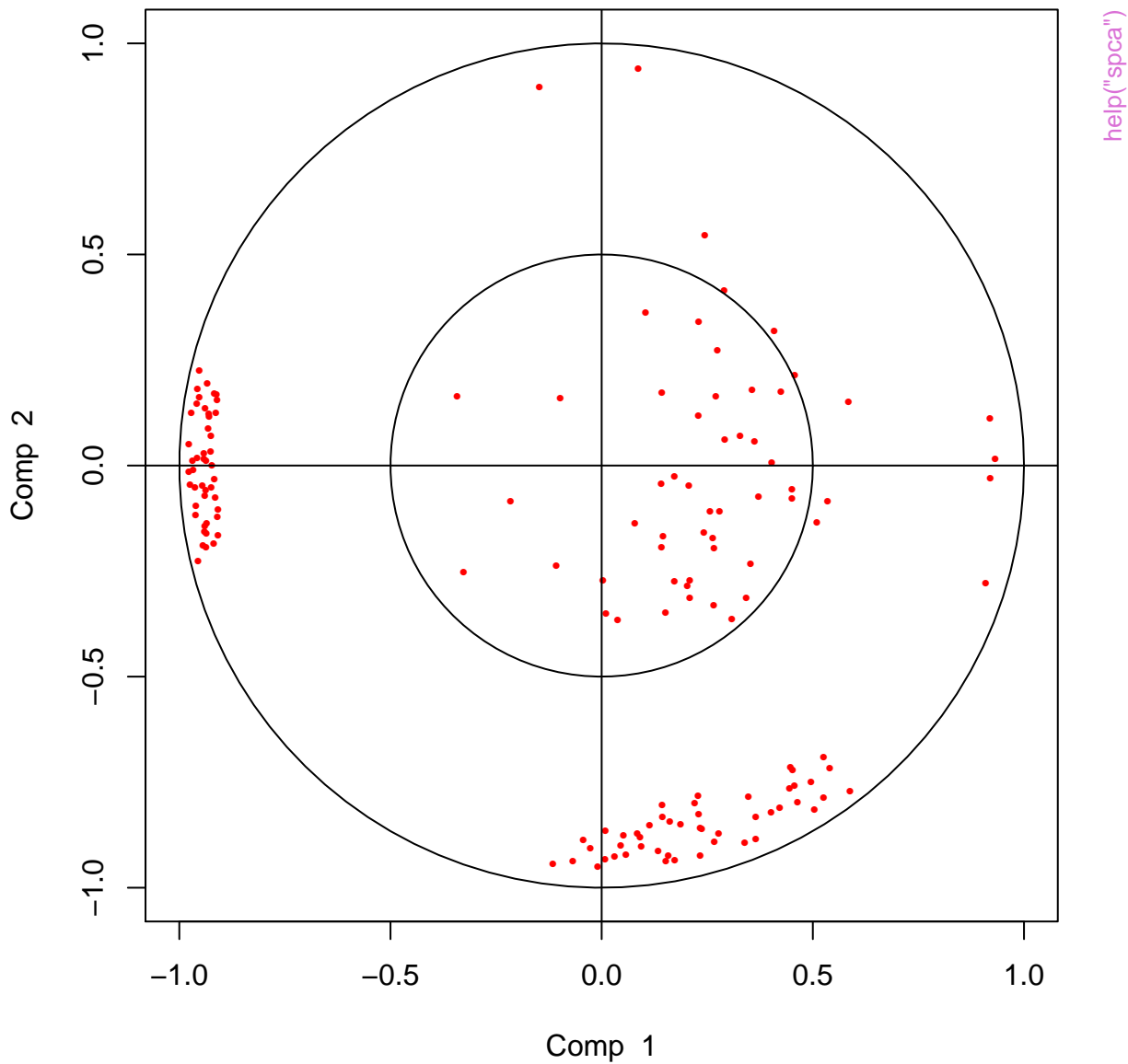


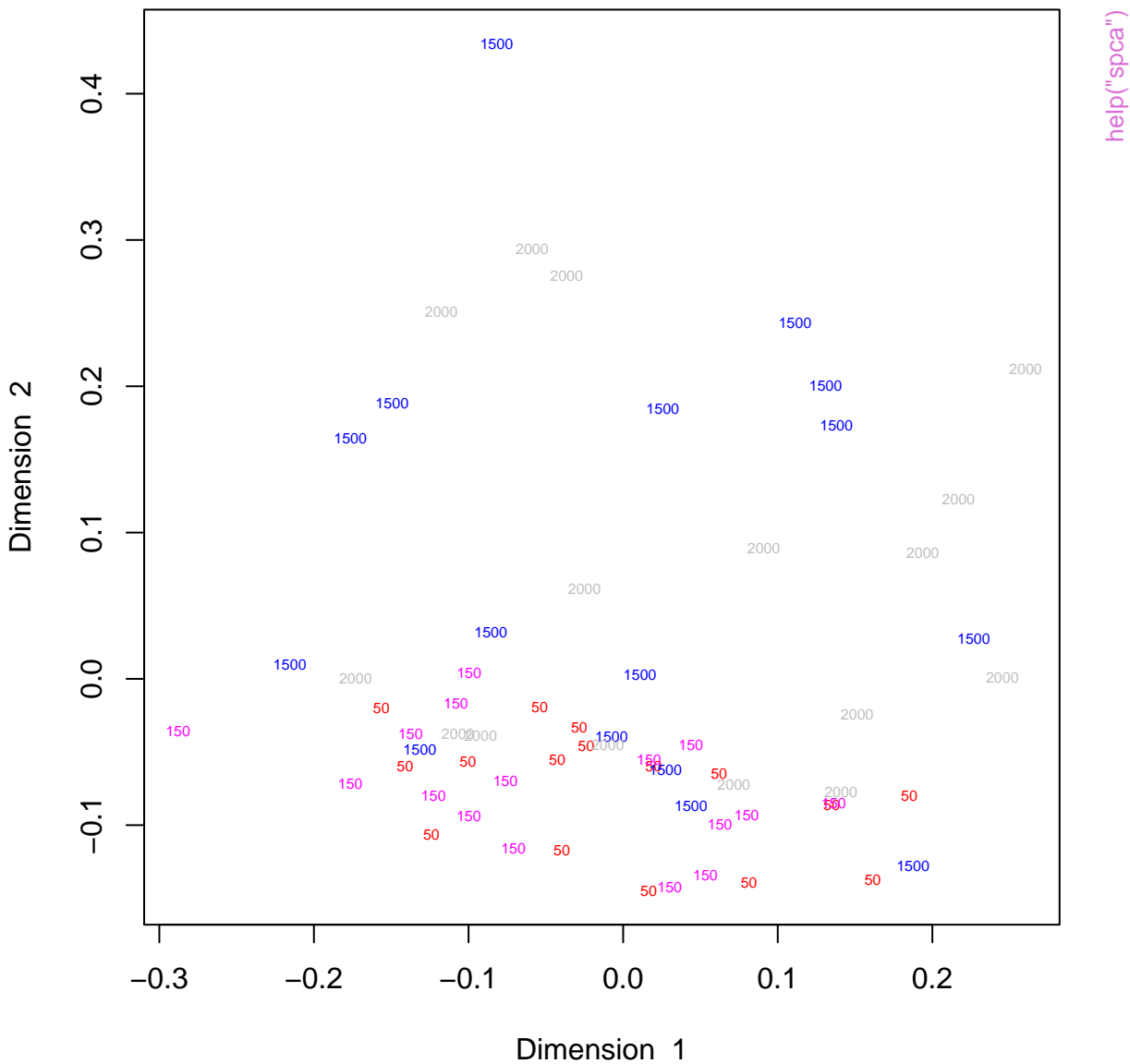


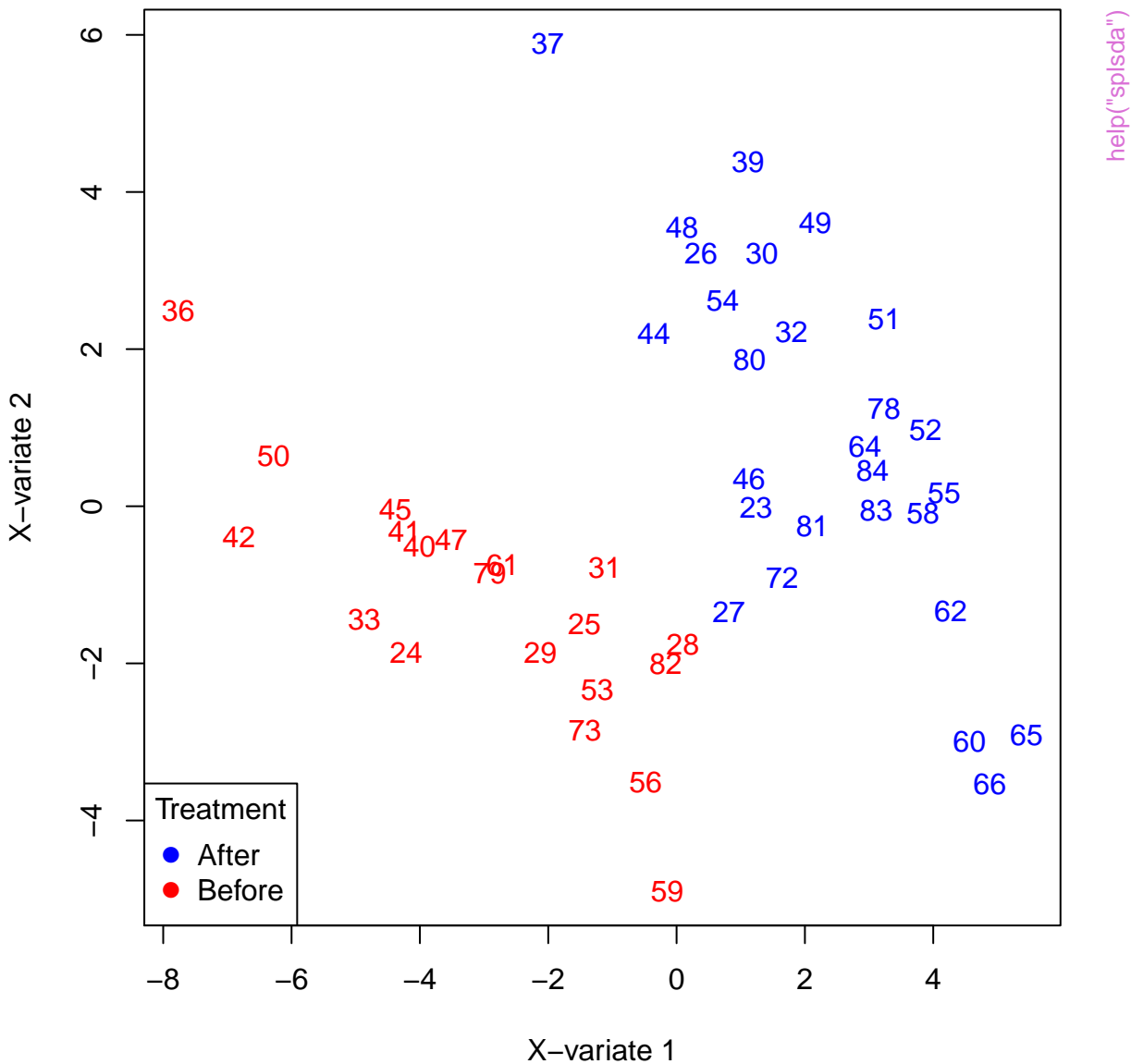
help("s.match")

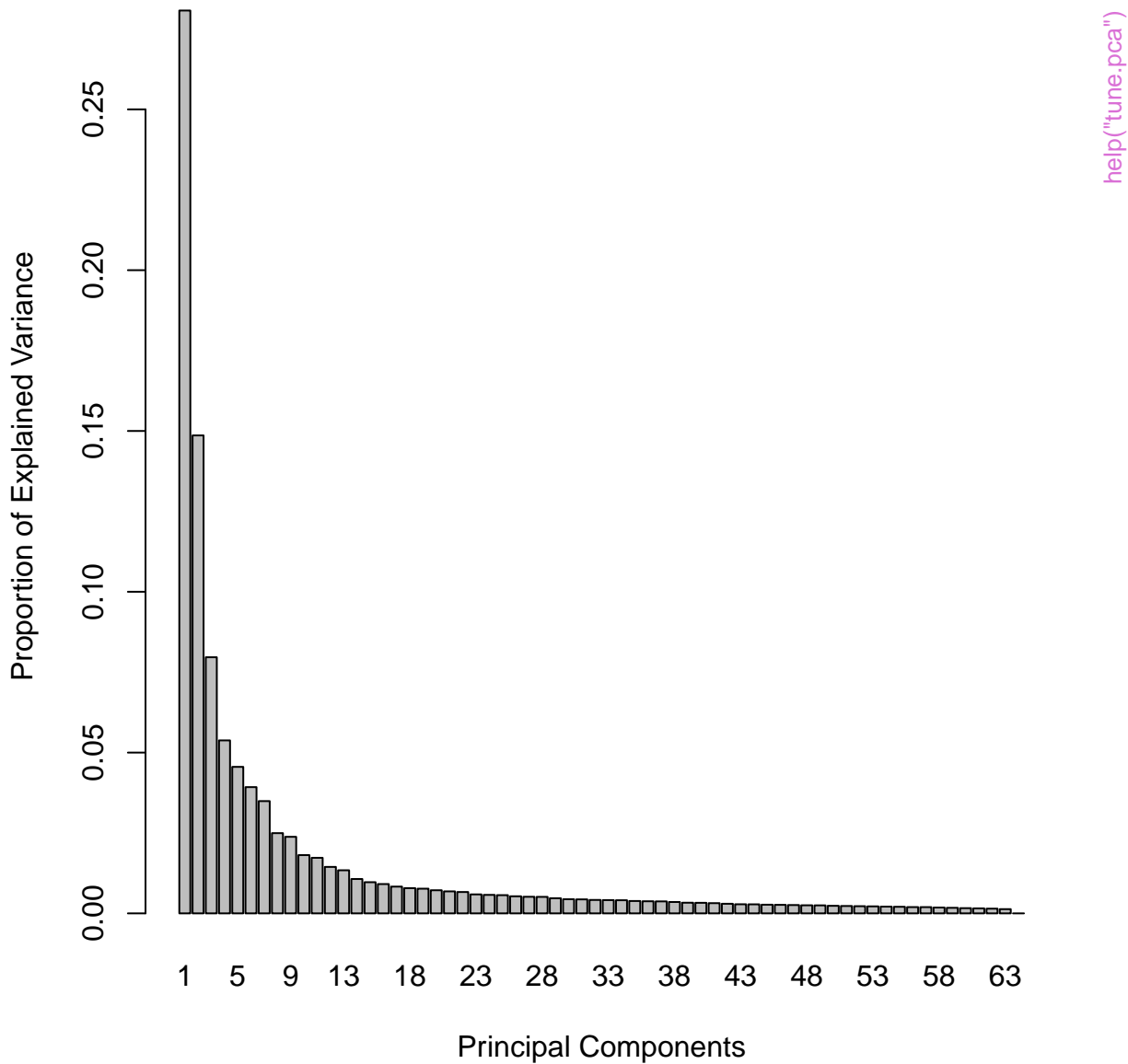












## ***Variable Importance in the Projection***

