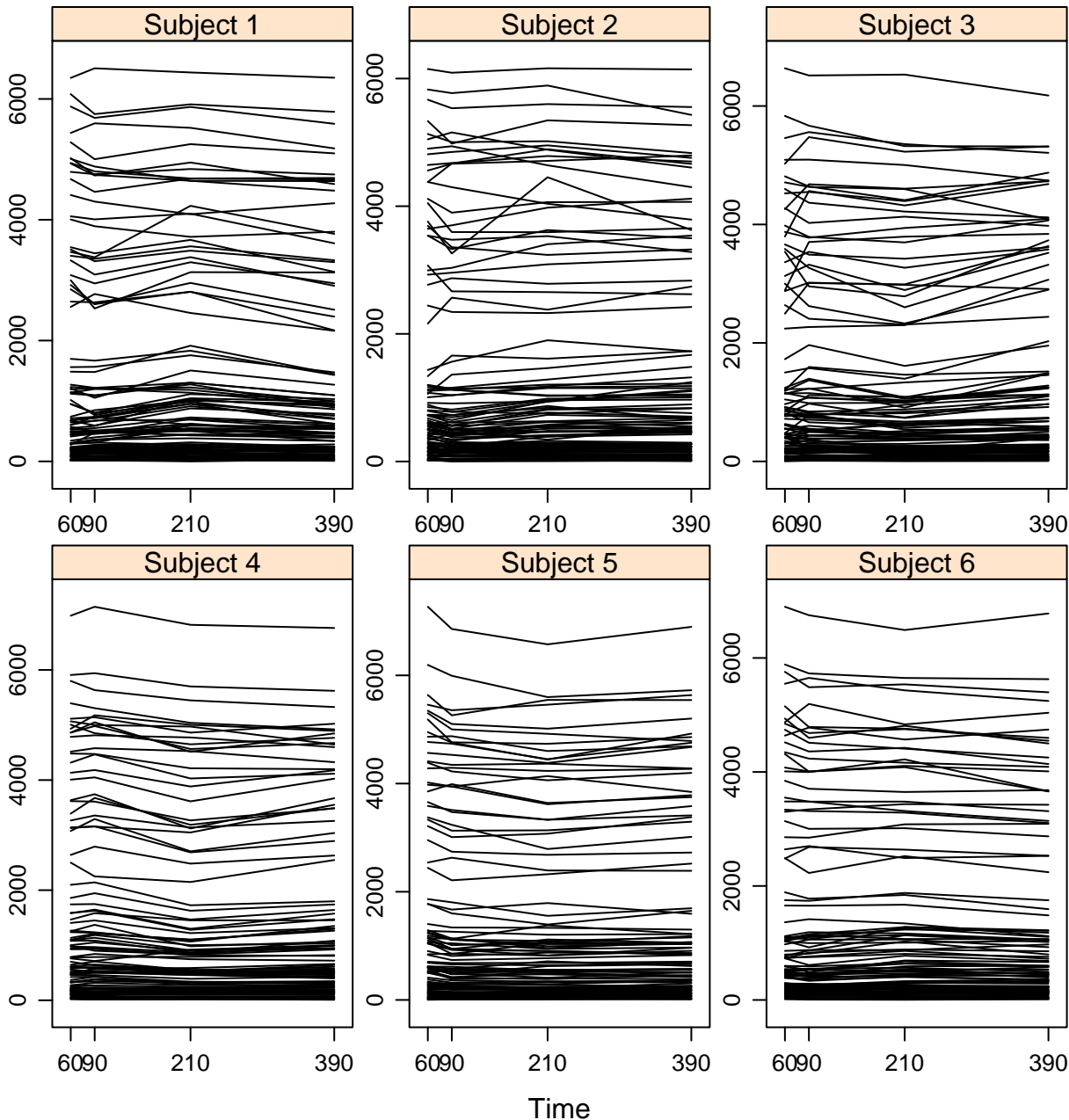
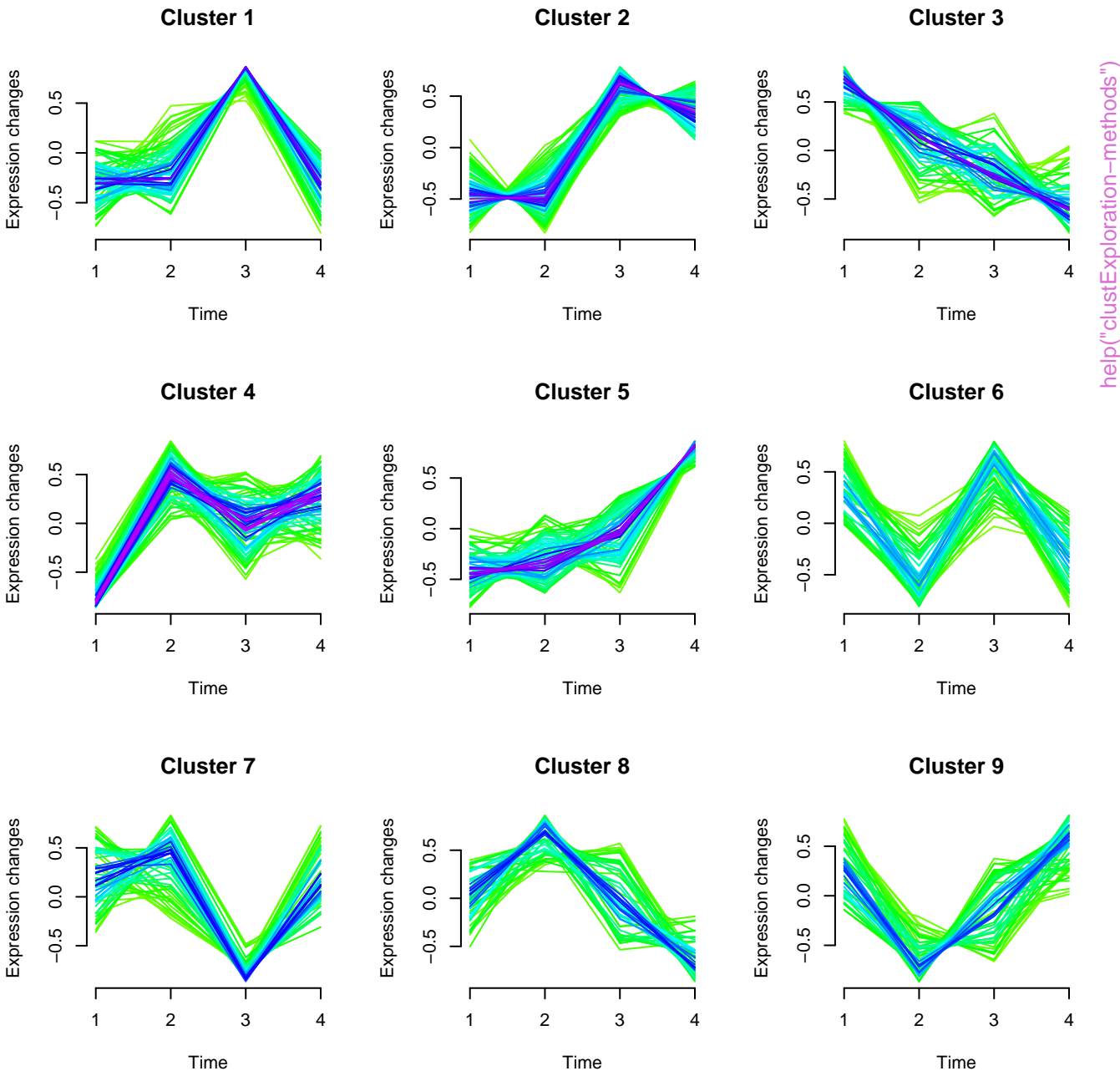
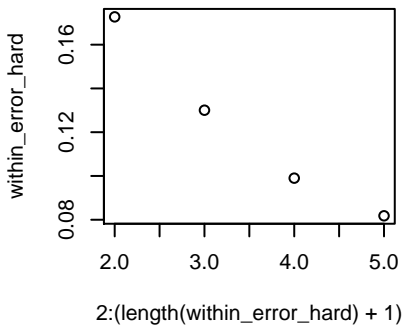


Gene Expression

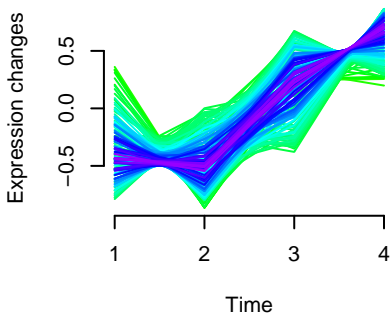


help("as.micro_array")

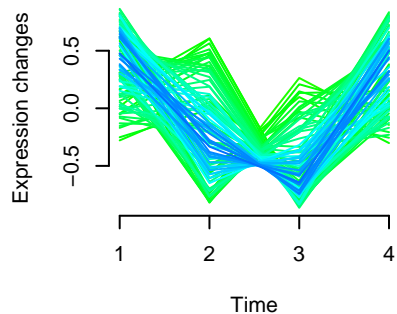




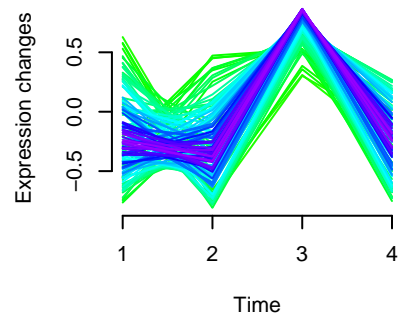
Cluster 1



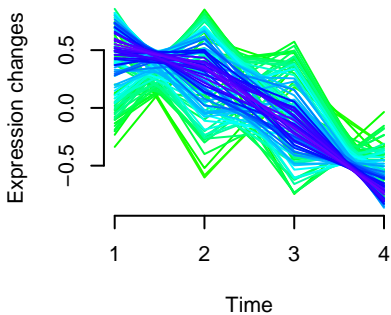
Cluster 2



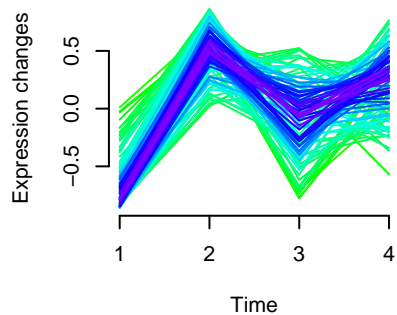
Cluster 3

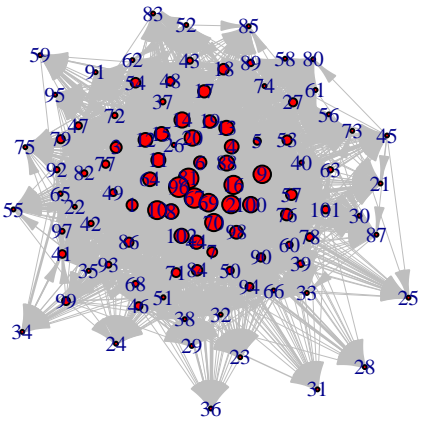


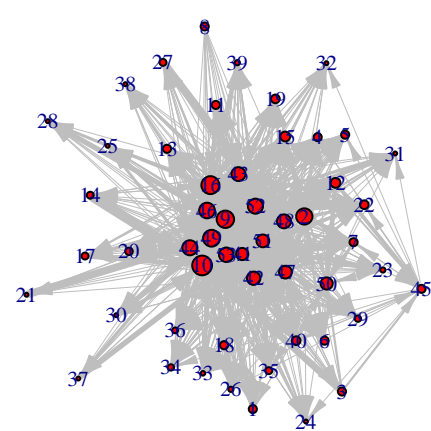
Cluster 4

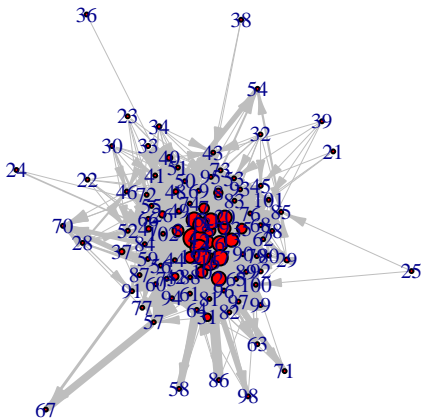


Cluster 5

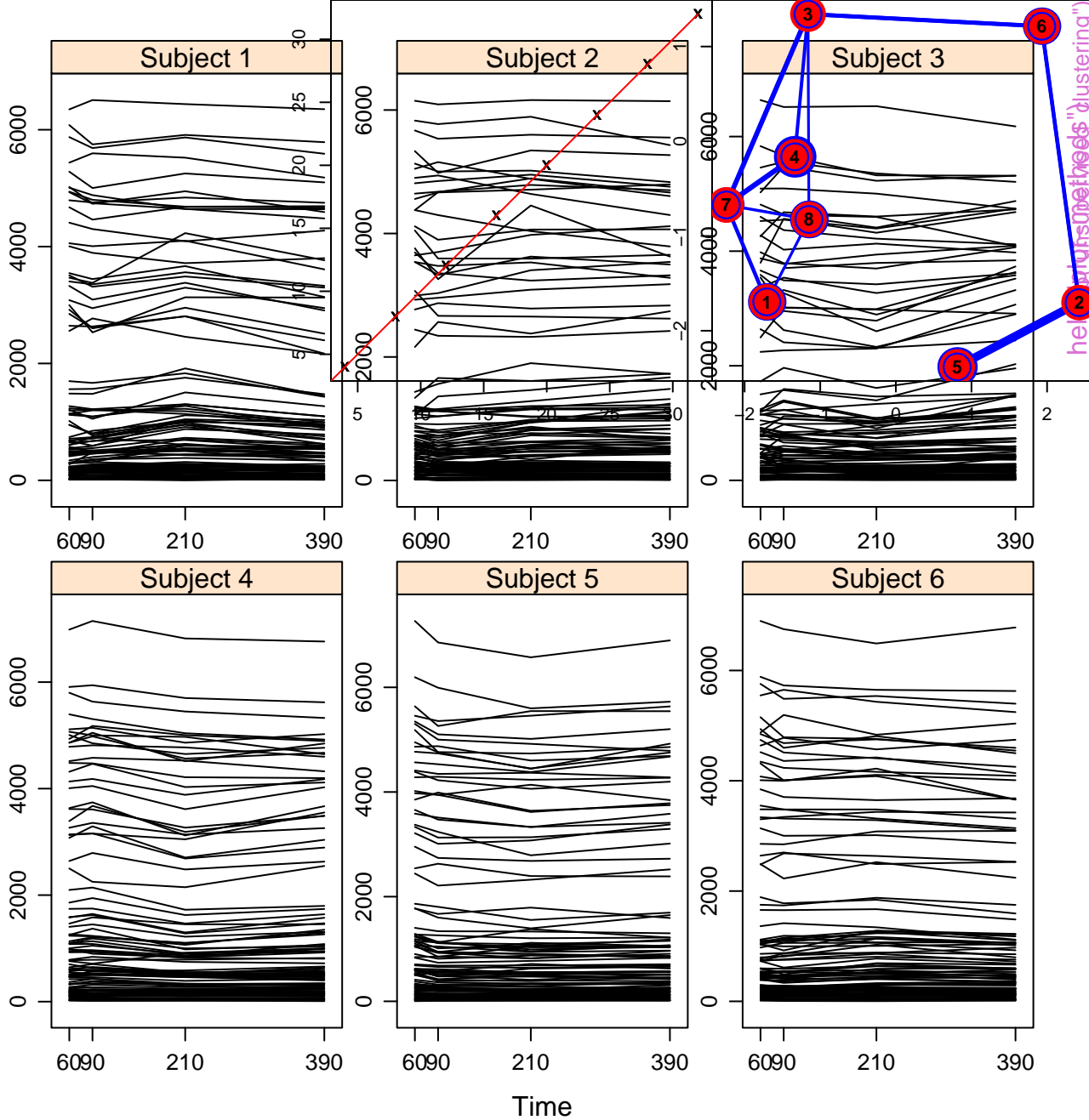




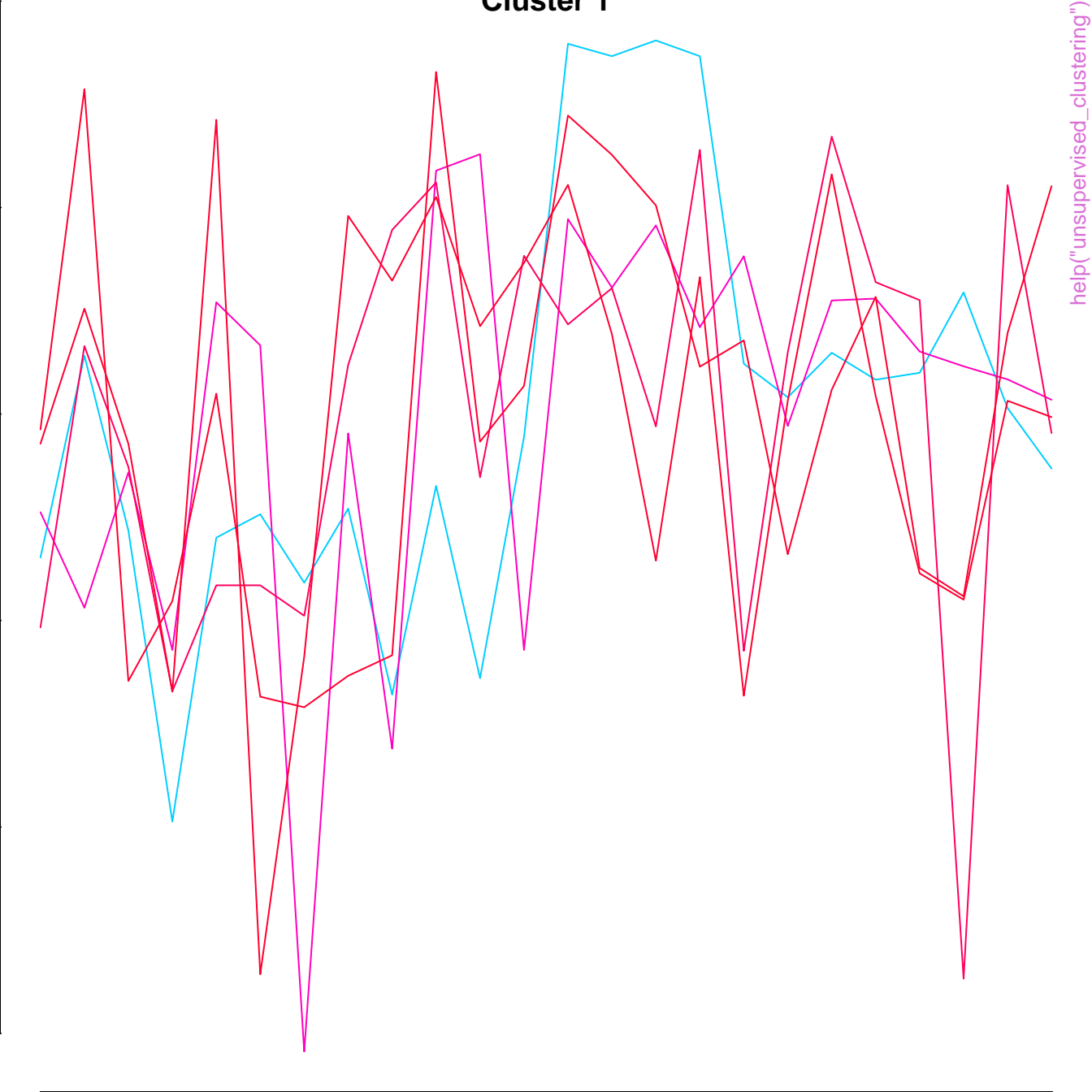




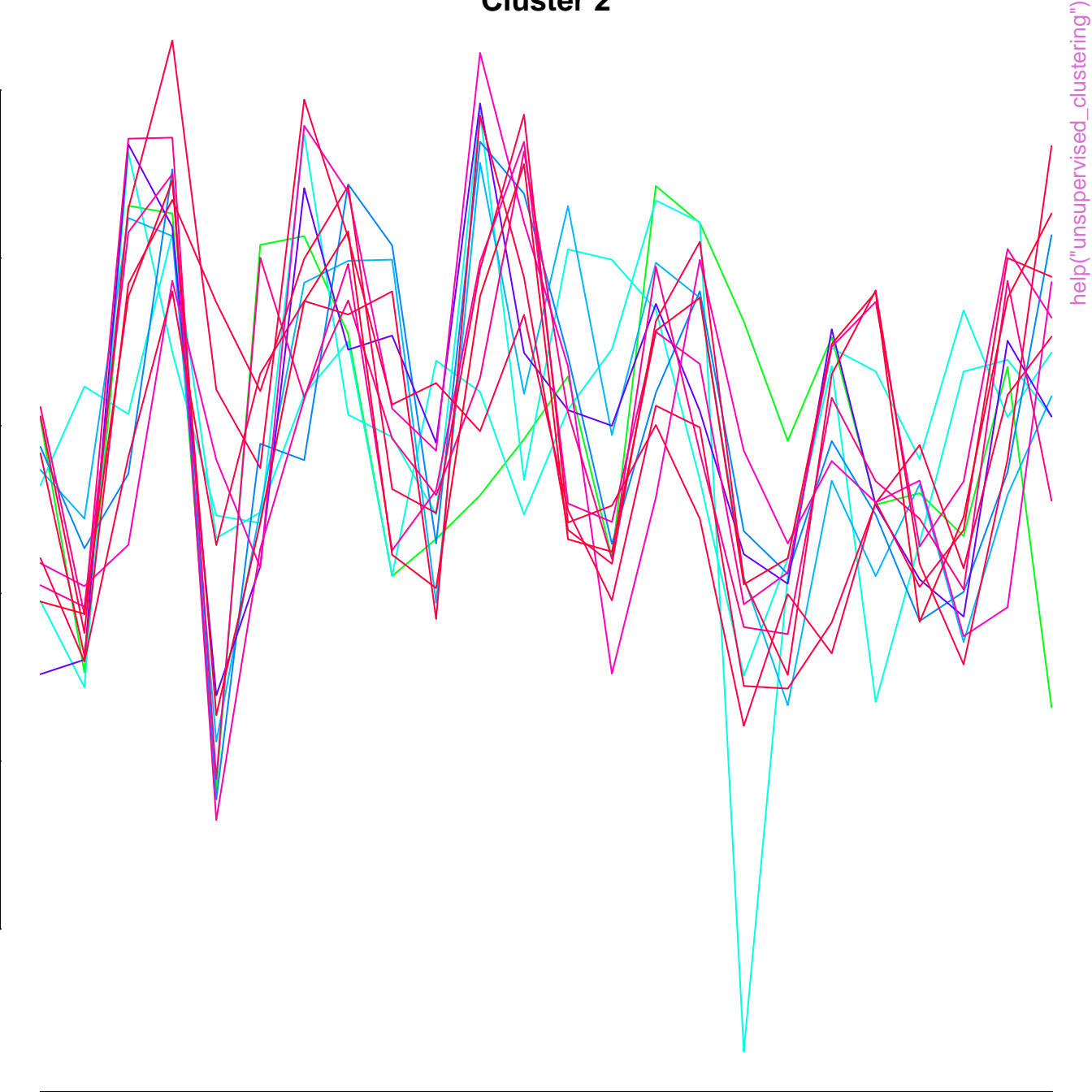
Gene Expression



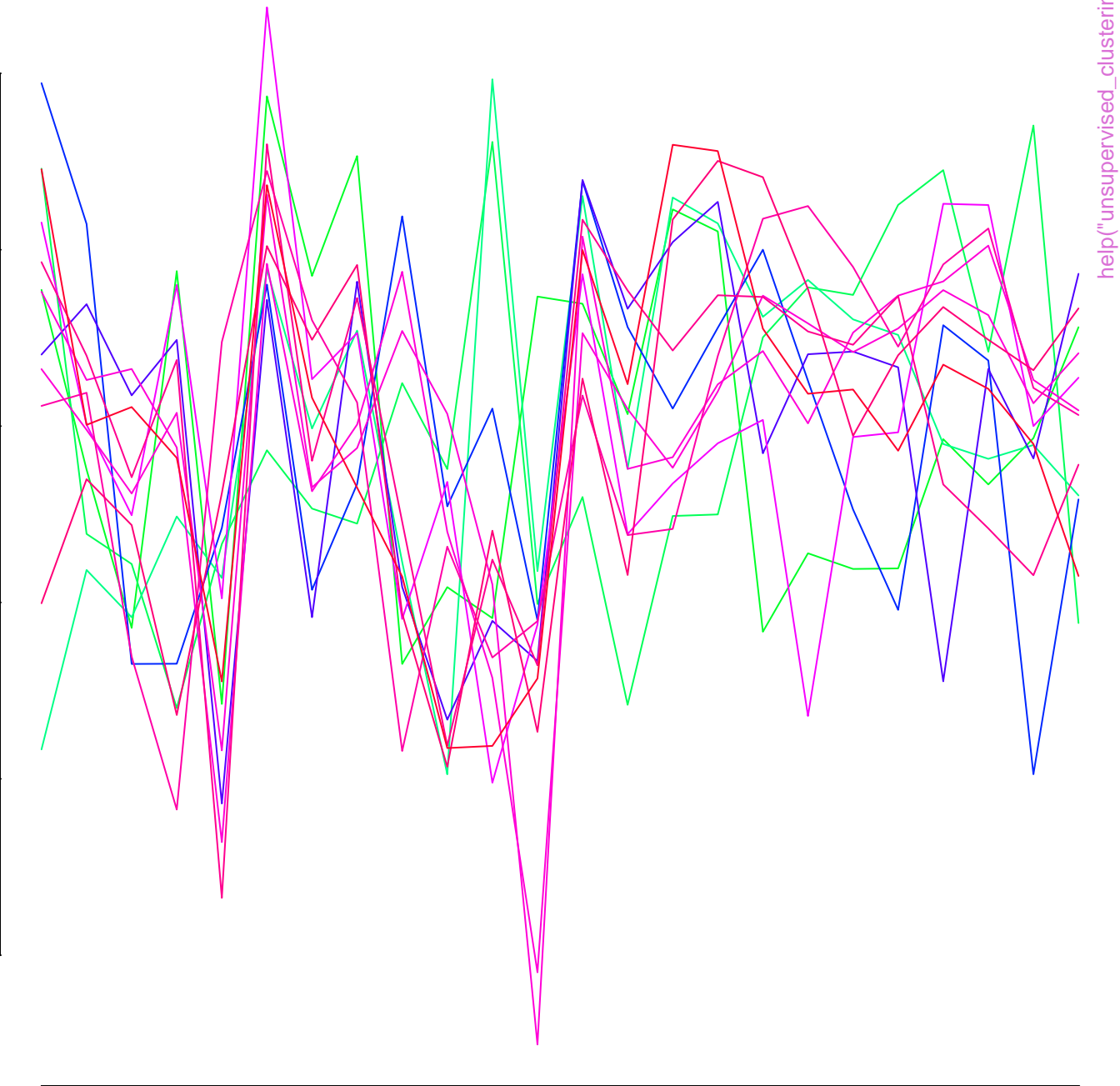
Cluster 1



Cluster 2

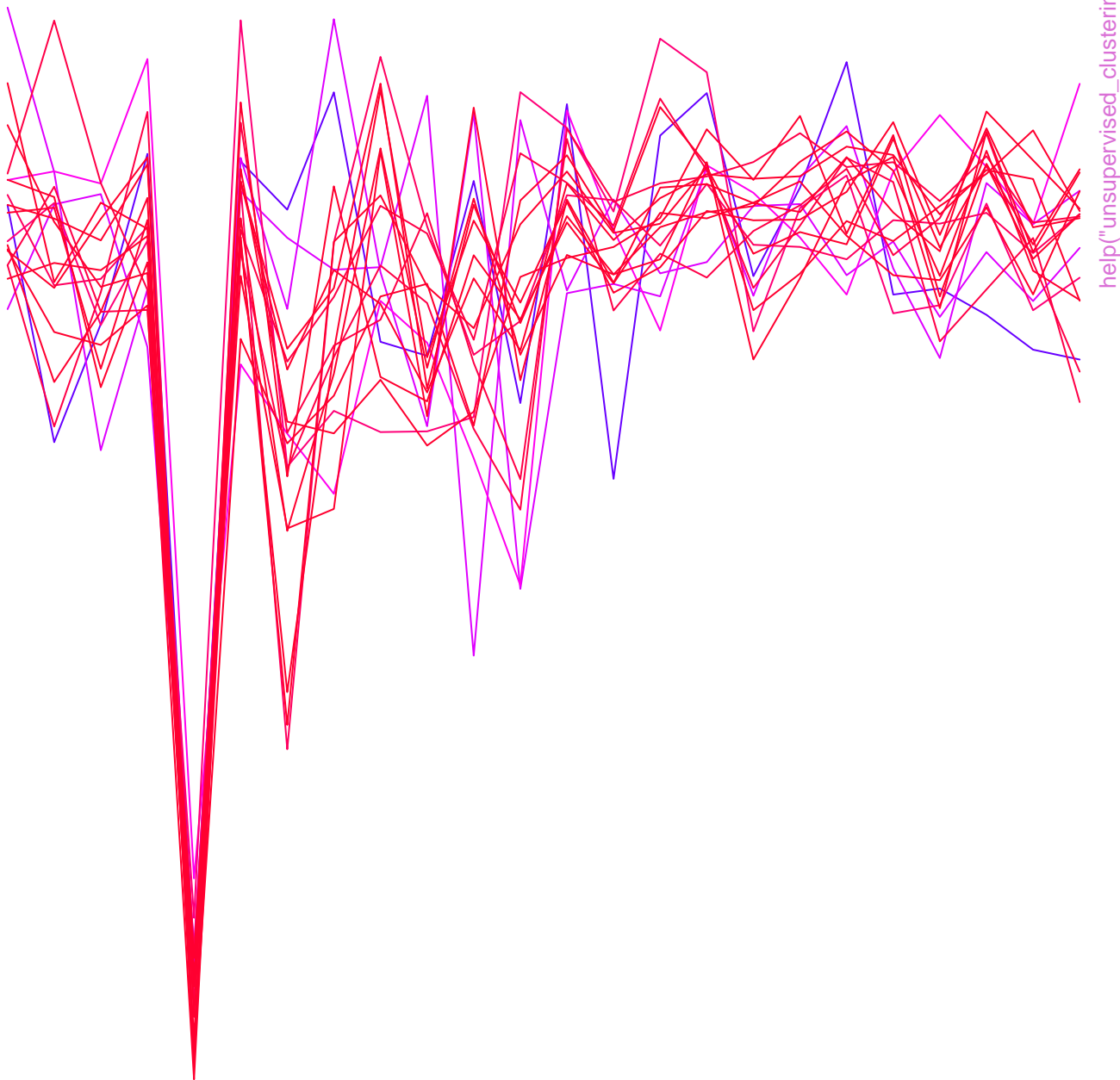


Cluster 3

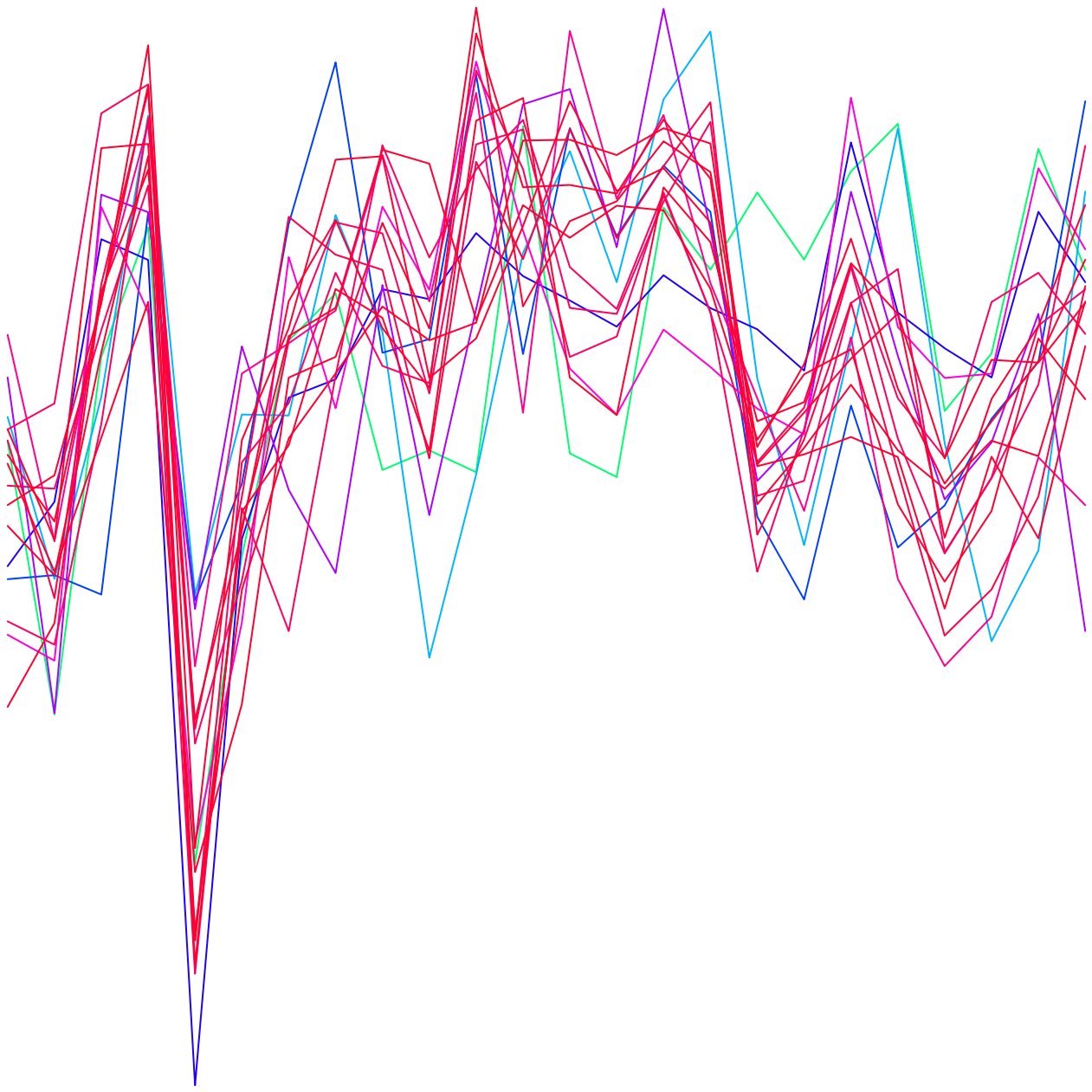


help("unsupervised_clustering")

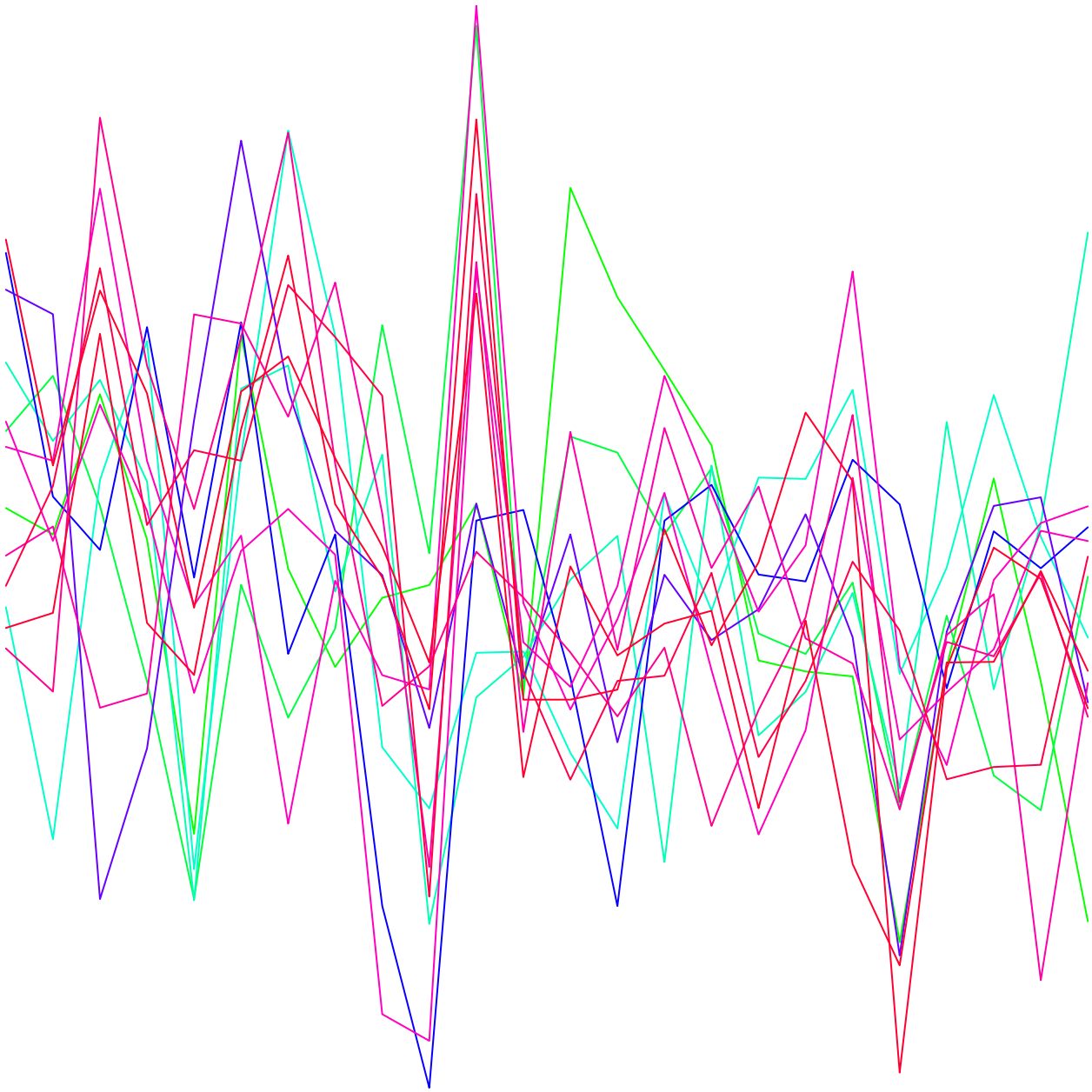
Cluster 4



Cluster 3

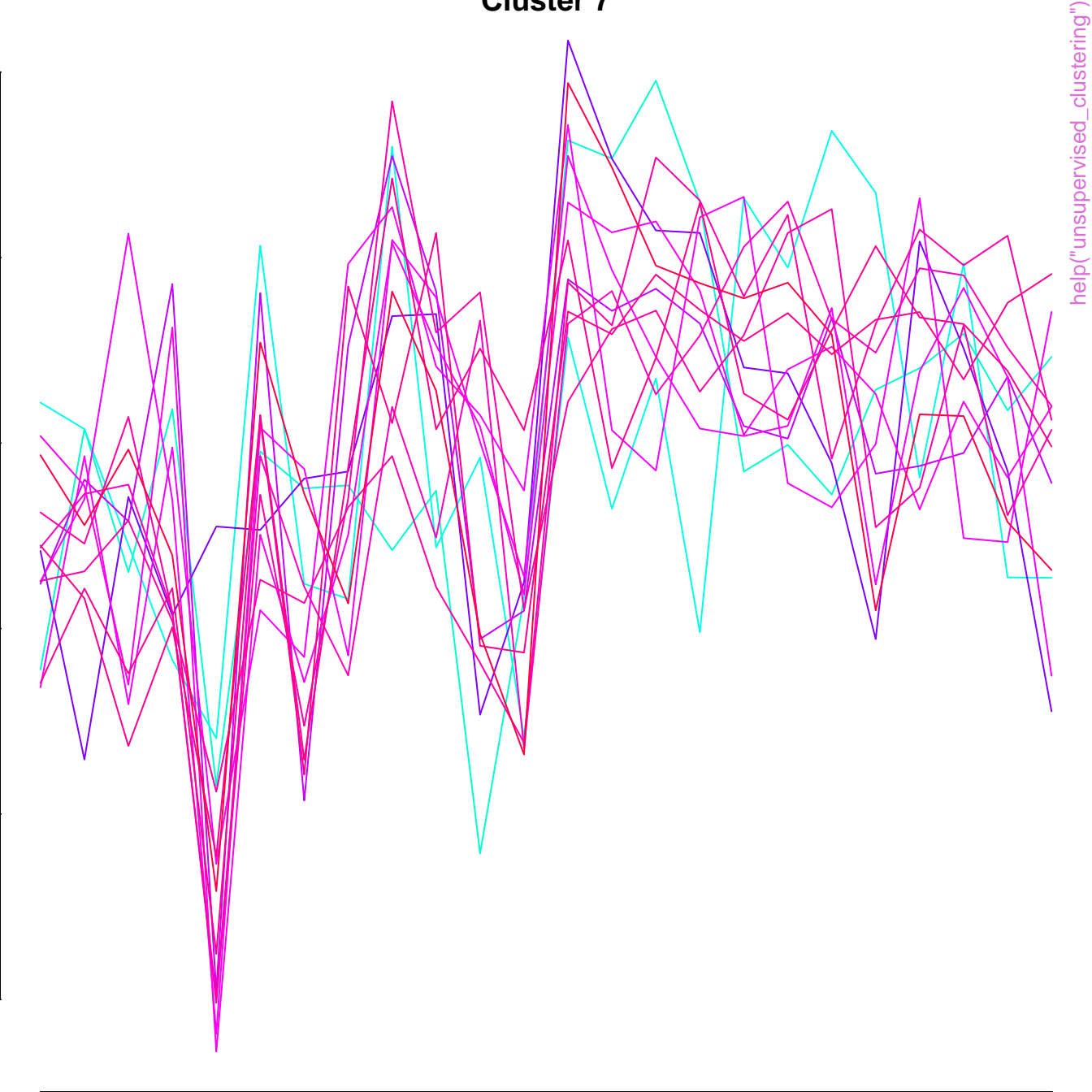


Cluster 6

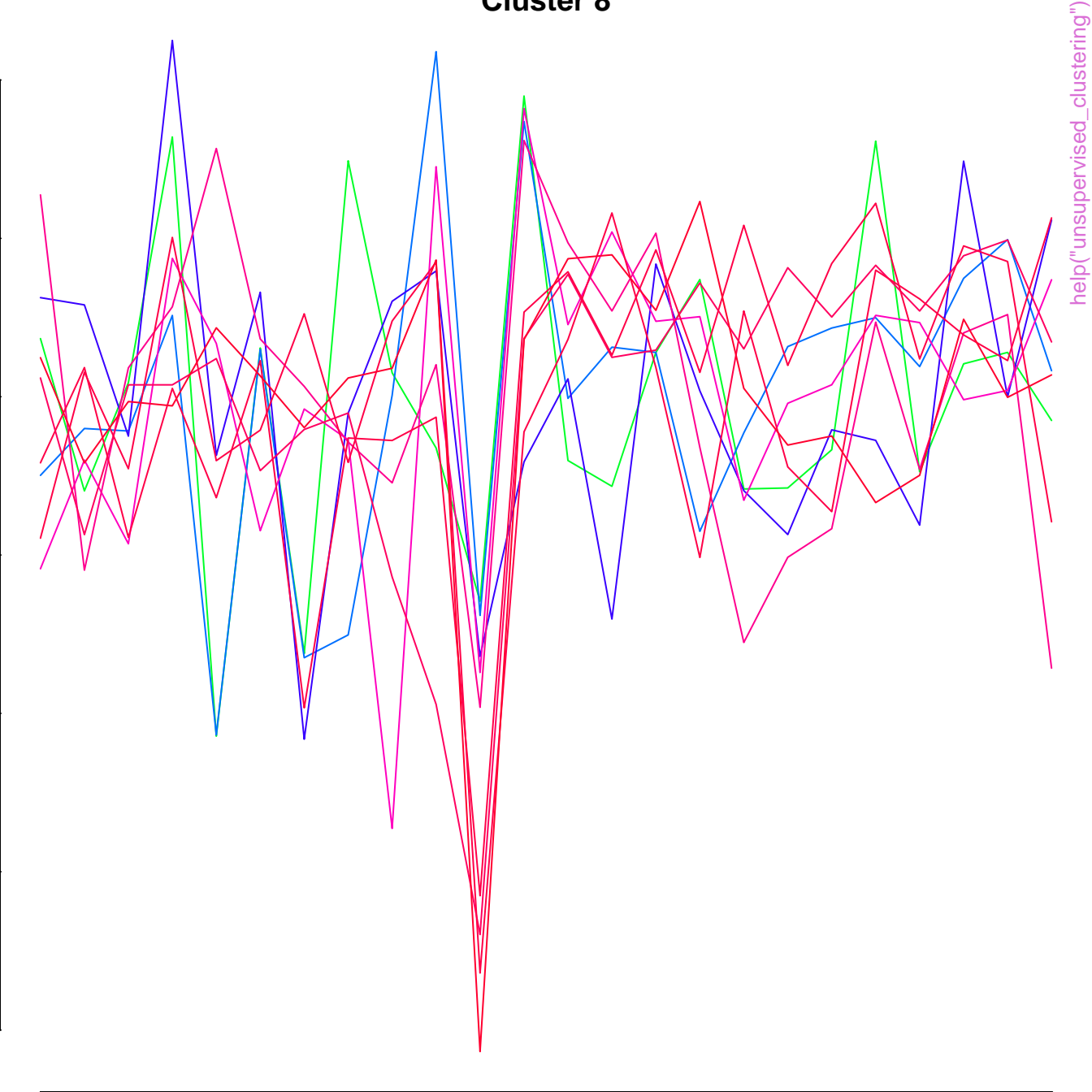


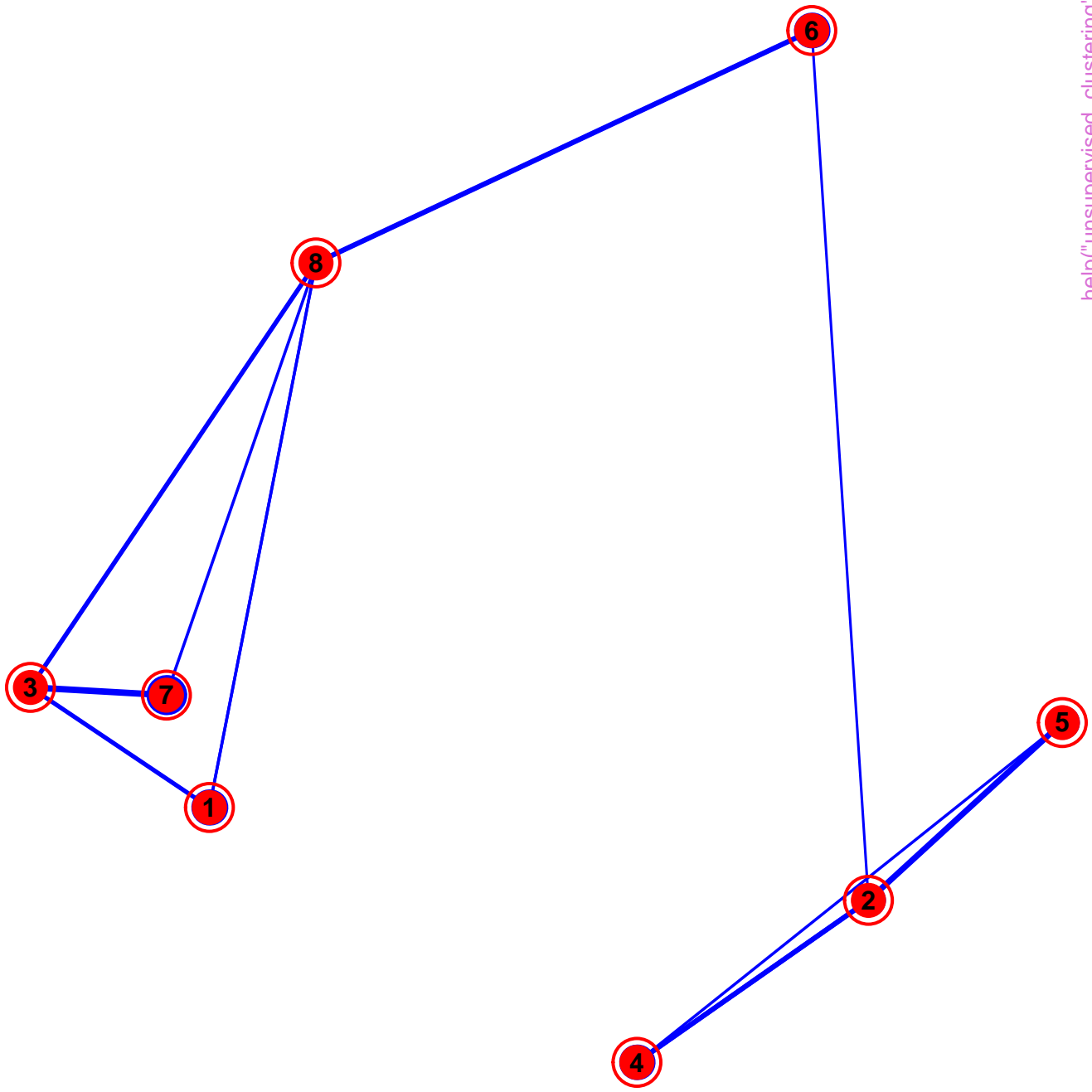
help("unsupervised_clustering")

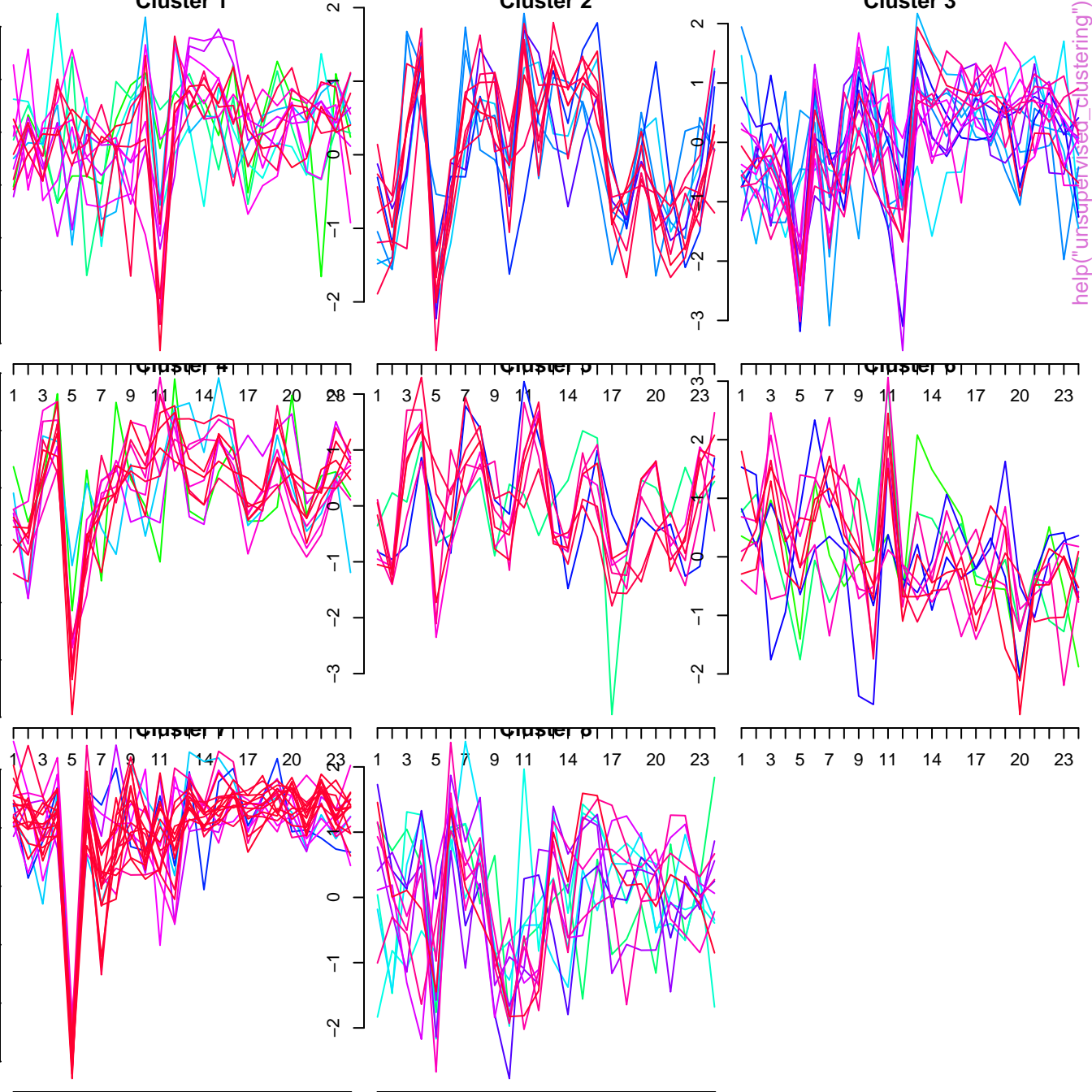
Cluster 7

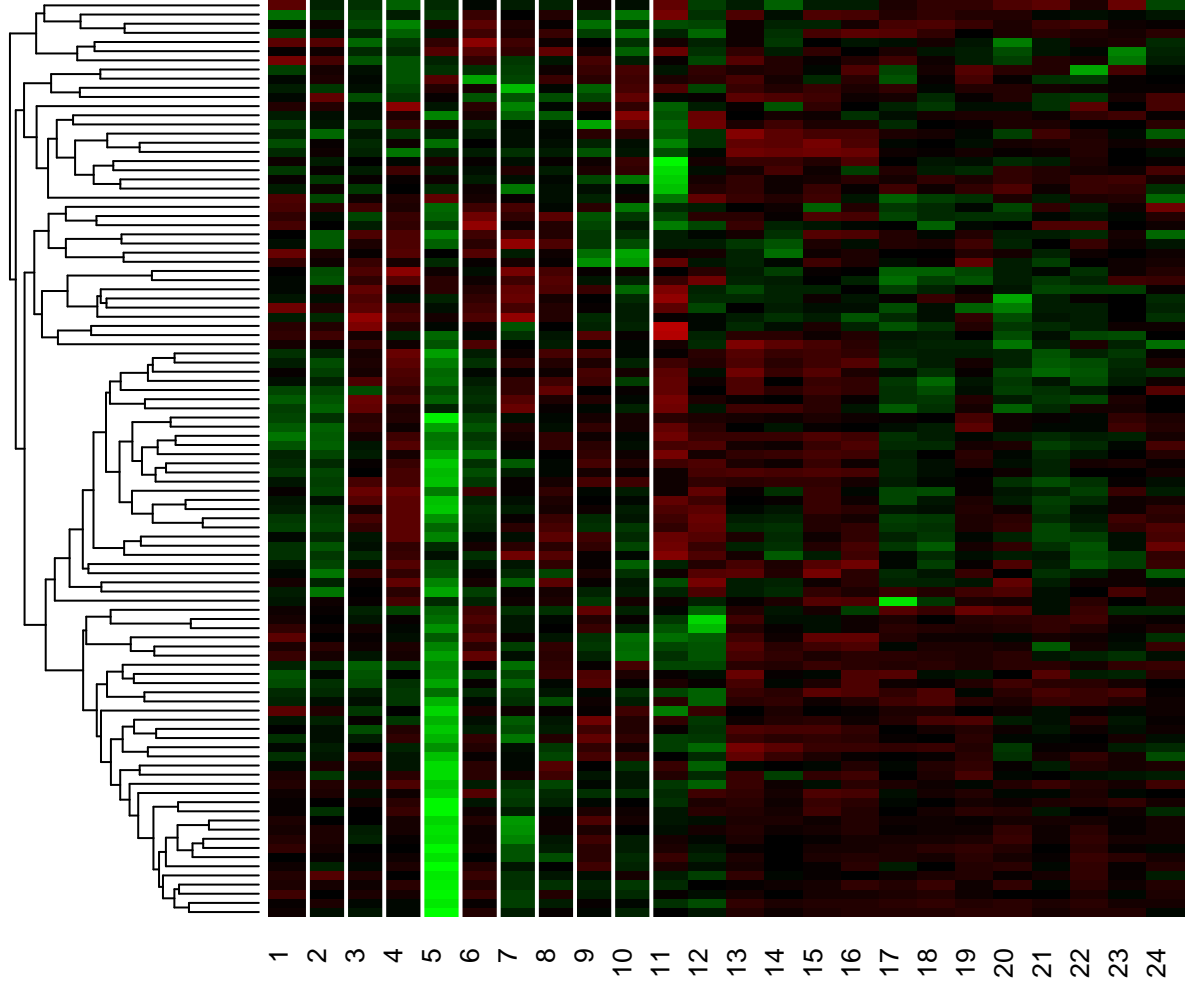


Cluster 8



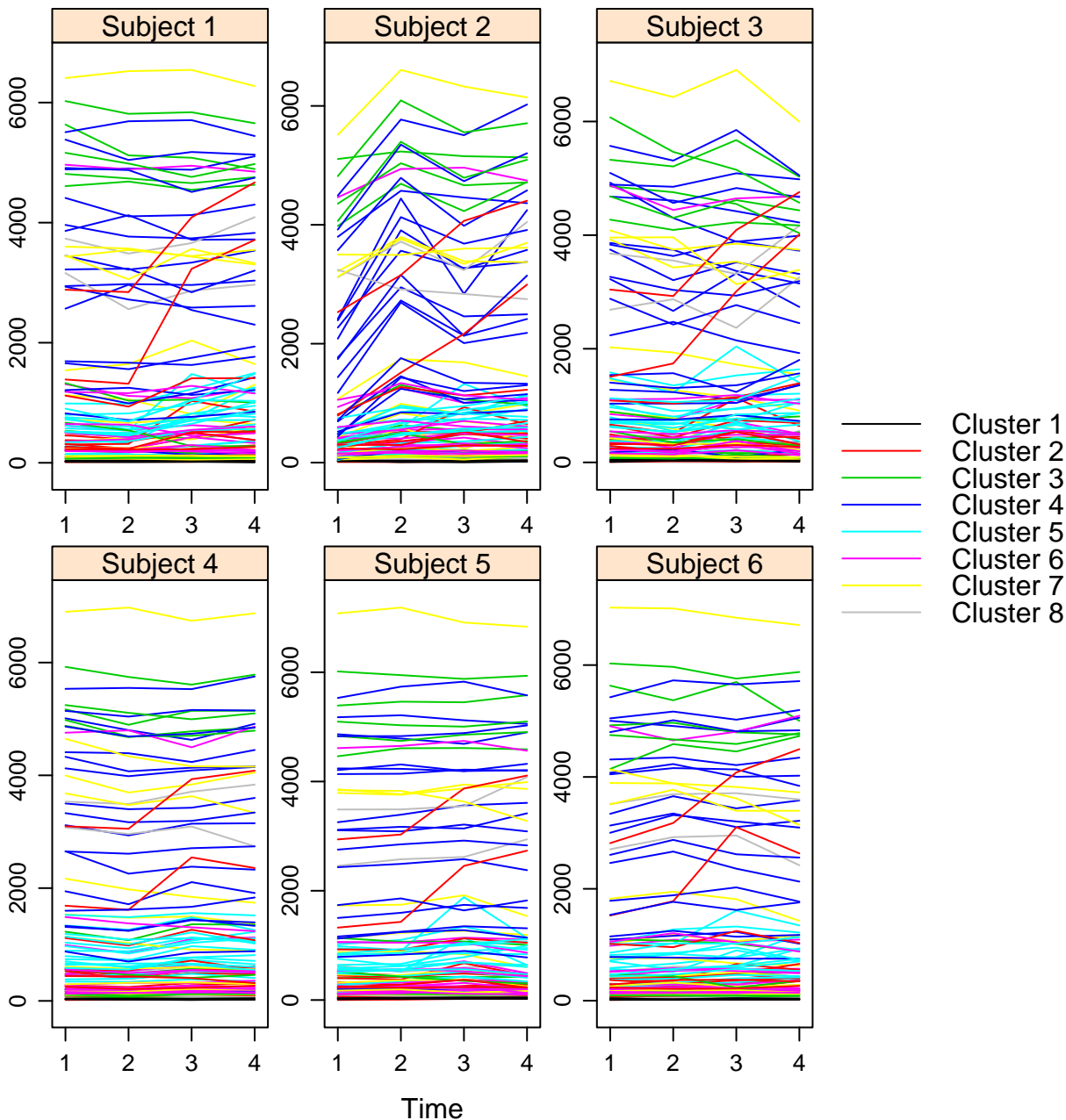






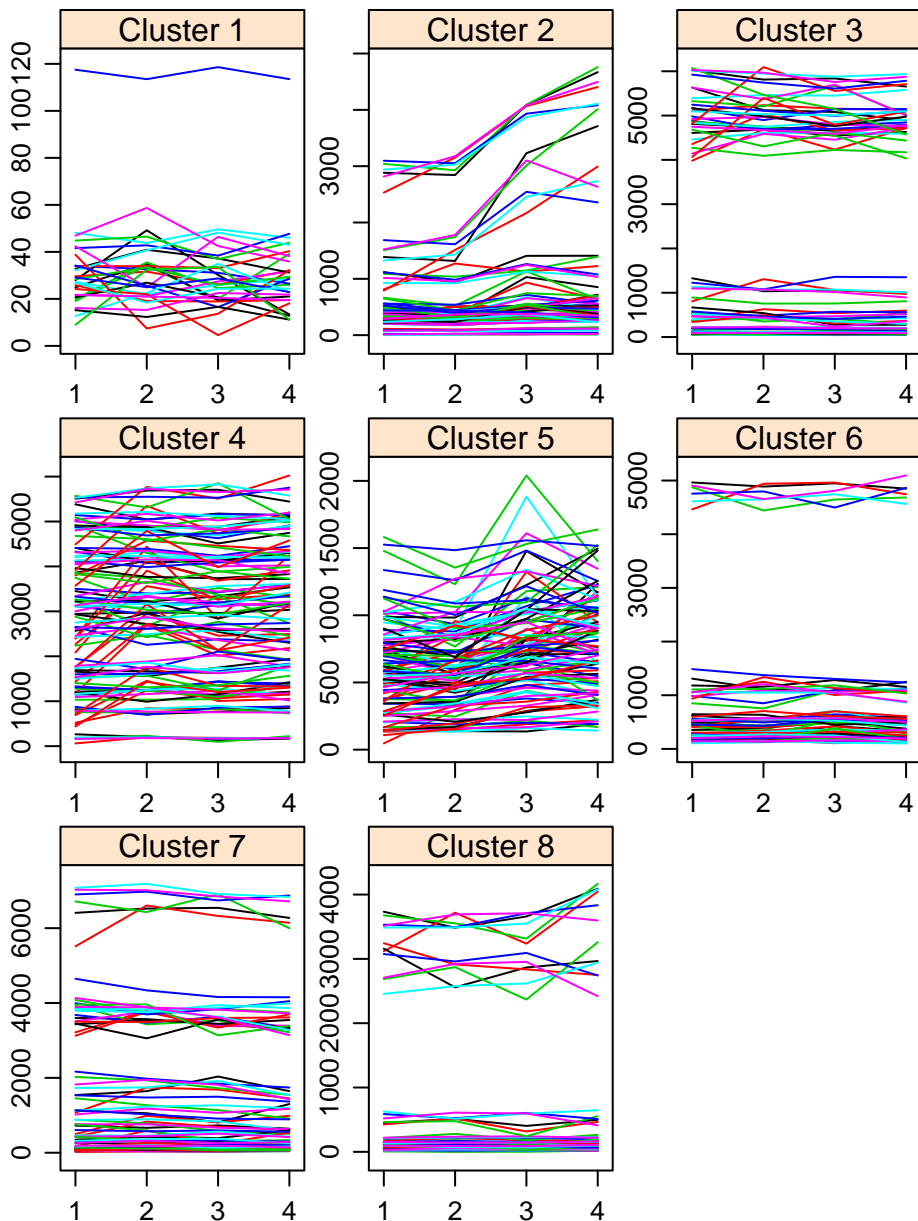
help("unsupervised_clustering")

Gene Expression



`help("unsupervised_clustering")`

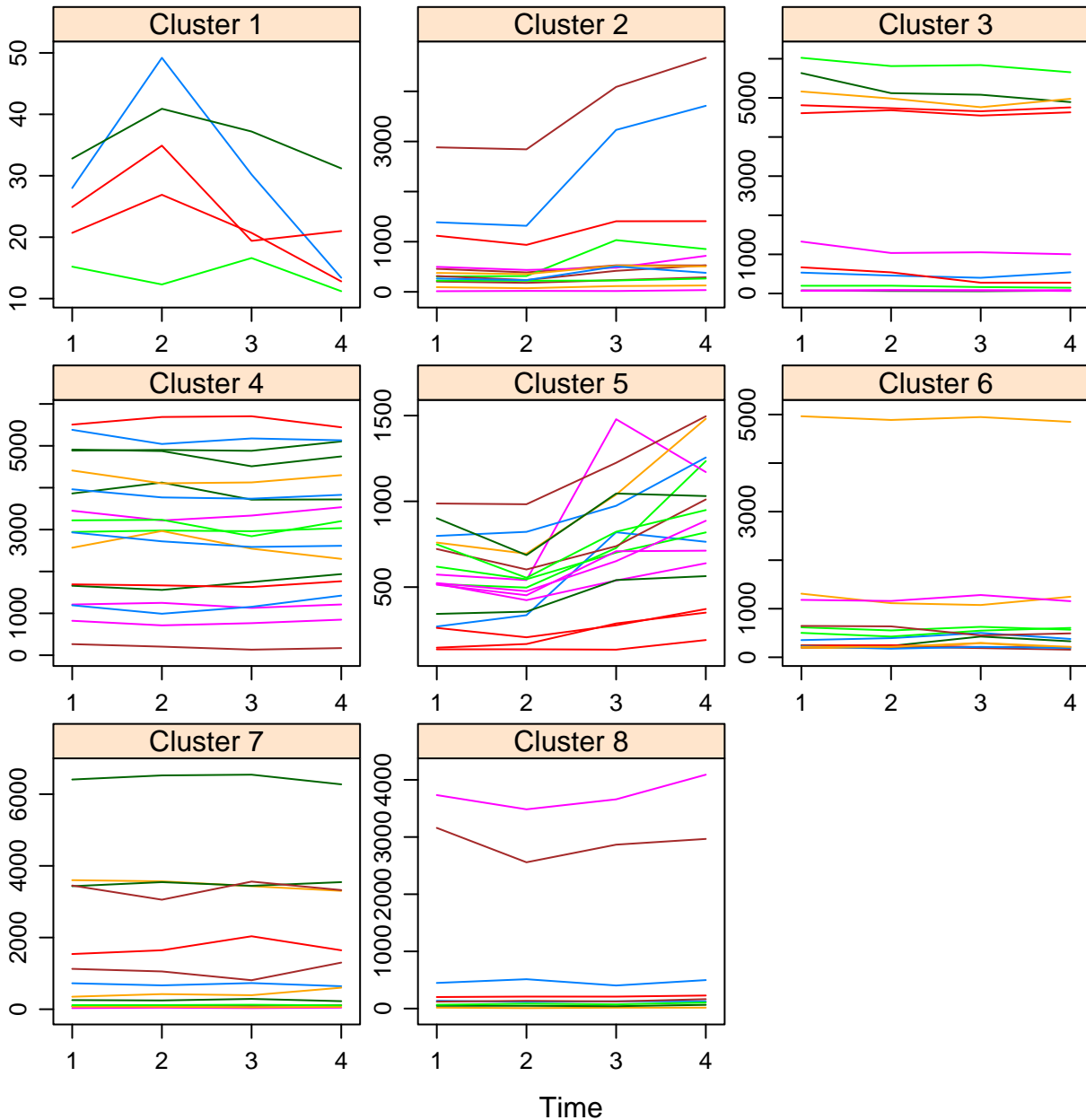
Gene Expression



— Subject 1
— Subject 2
— Subject 3
— Subject 4
— Subject 5
— Subject 6

help("unsupervised_clustering")

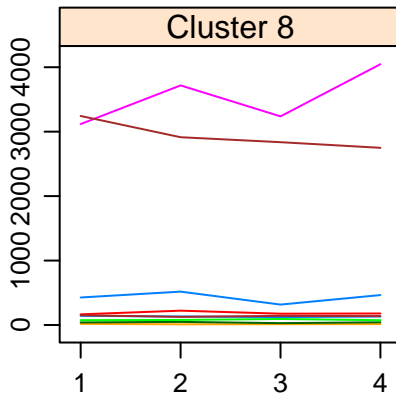
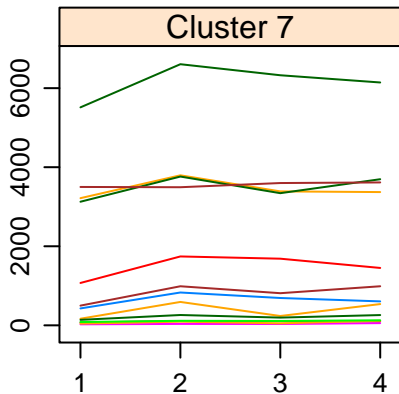
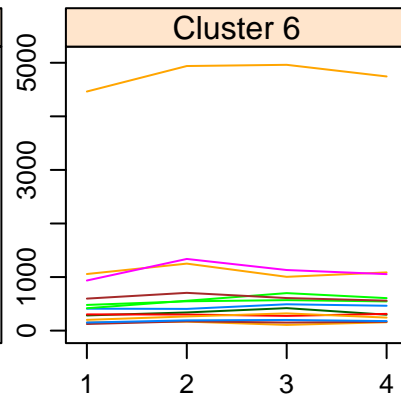
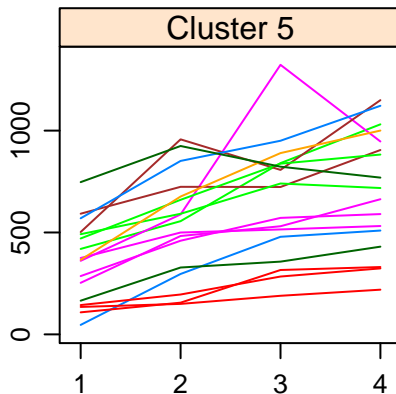
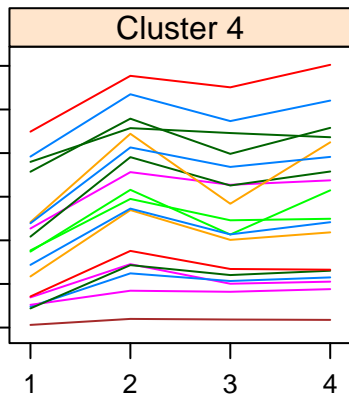
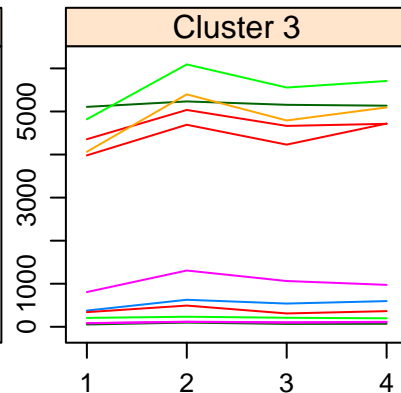
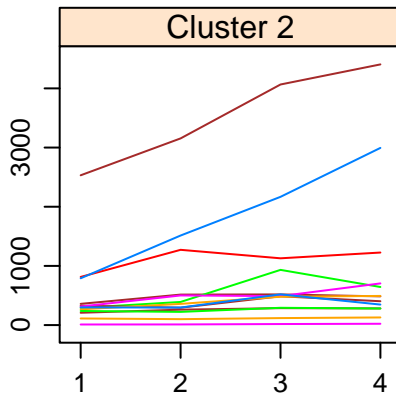
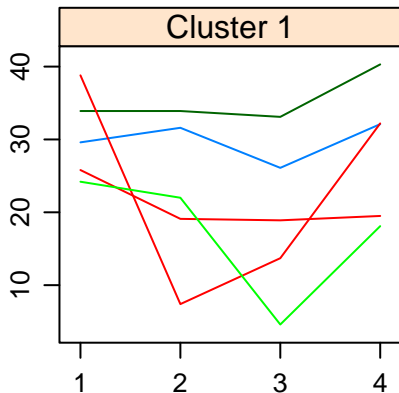
Subject 1



help("unsupervised_clustering")

Subject 2

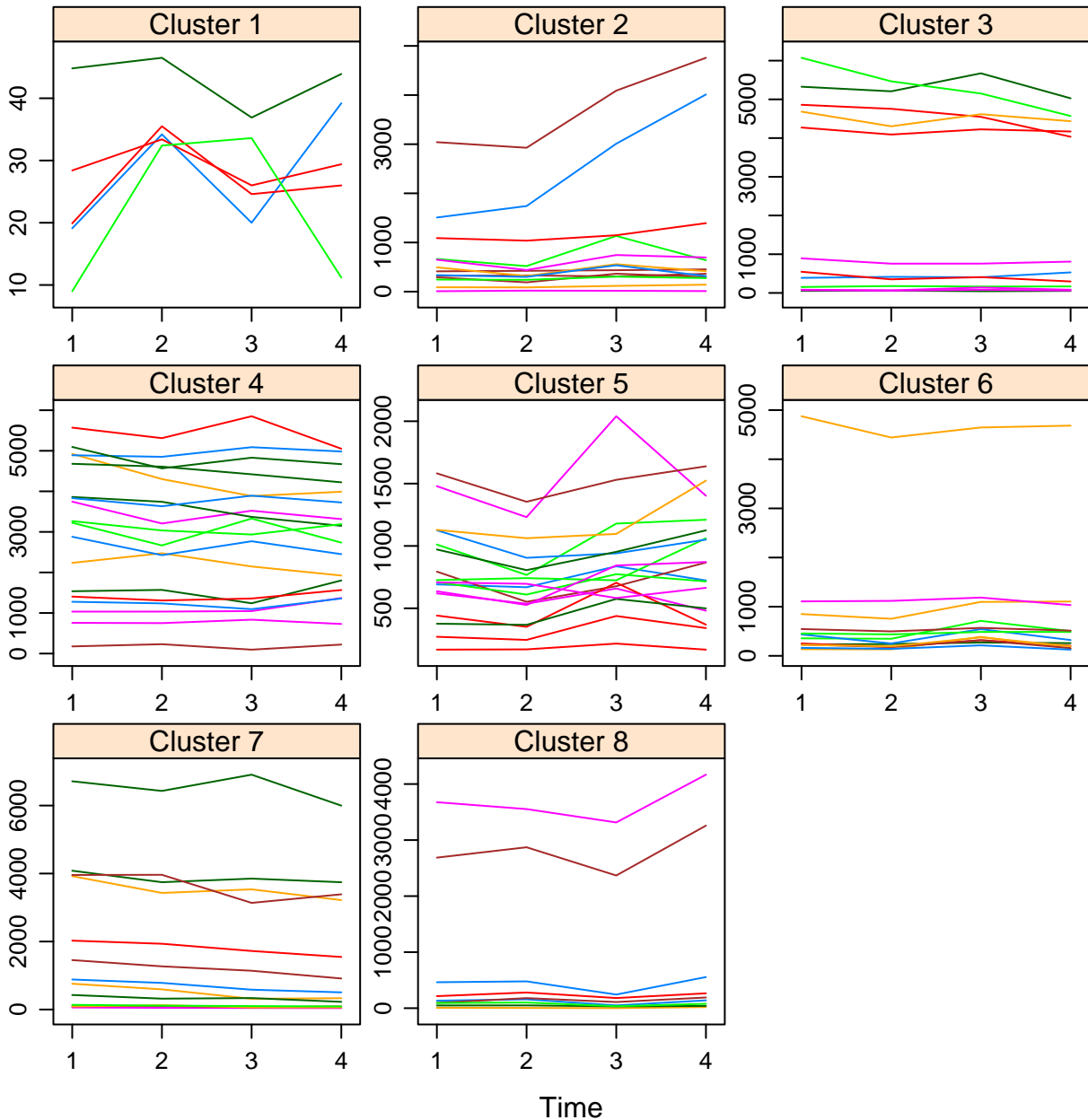
Gene Expression



Time

help("unsupervised_clustering")

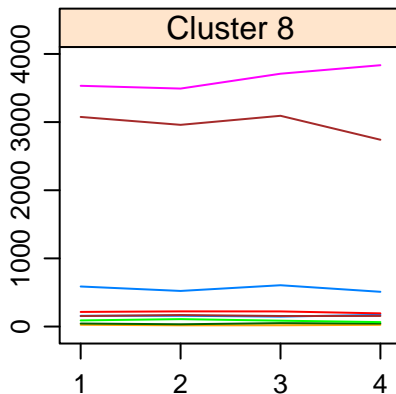
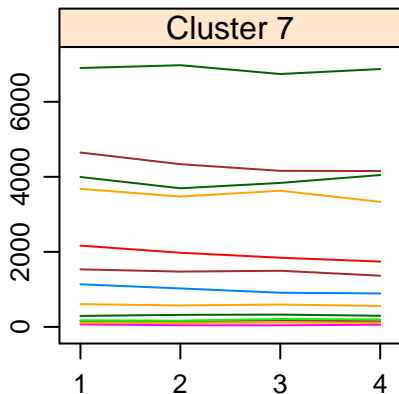
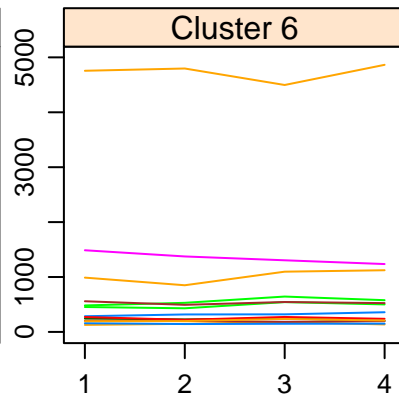
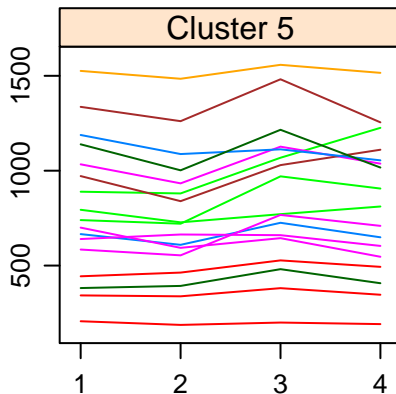
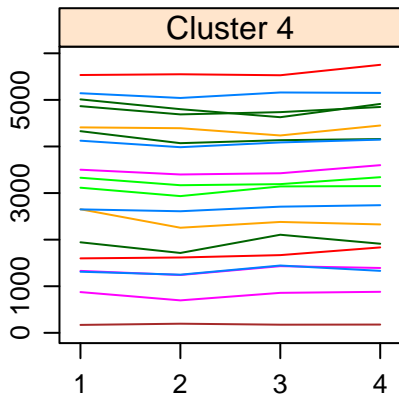
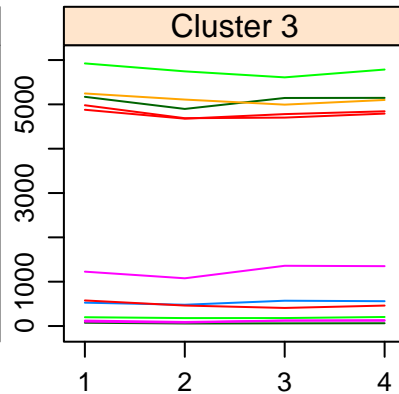
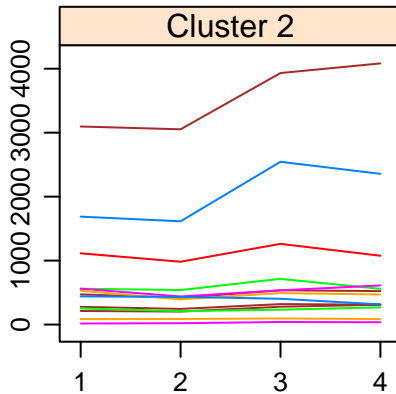
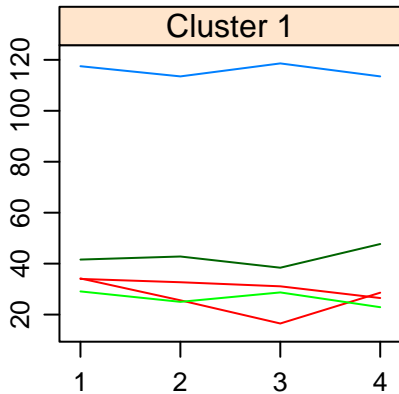
Subject 3



help("unsupervised_clustering")

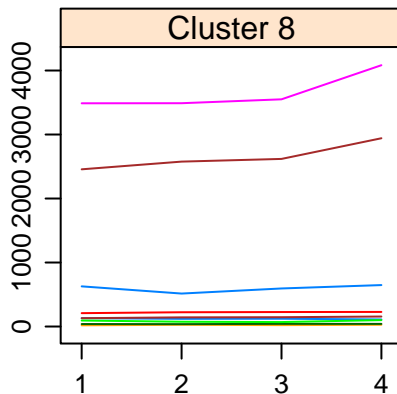
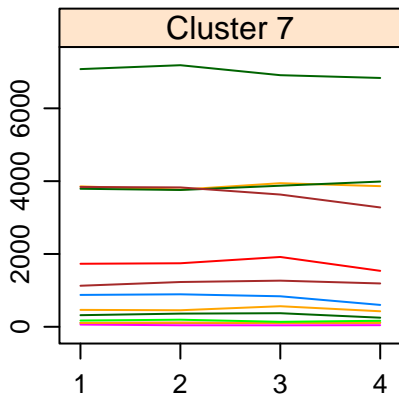
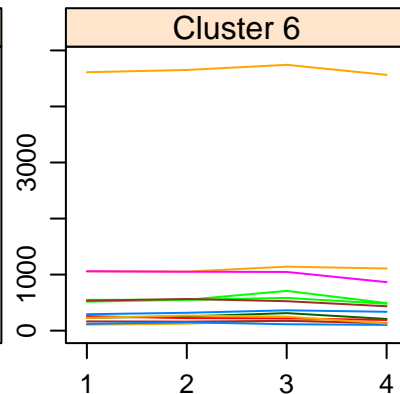
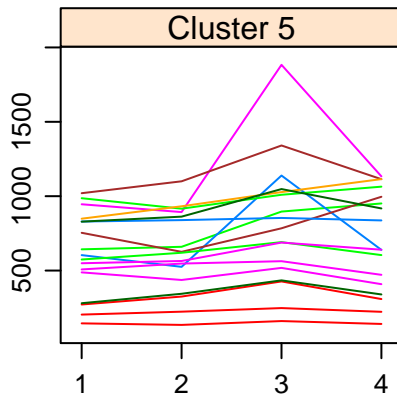
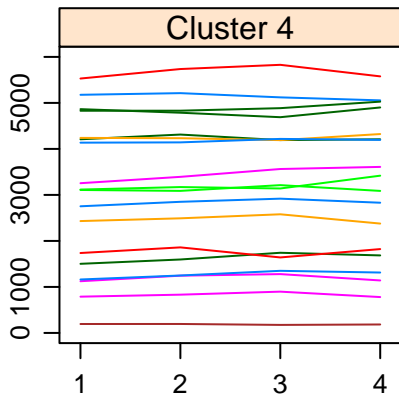
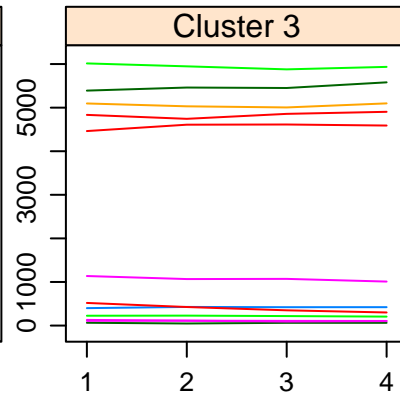
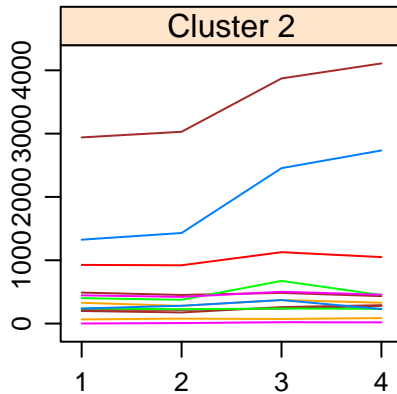
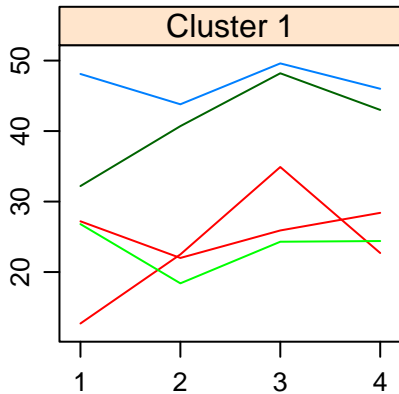
Subject 4

Gene Expression



help("unsupervised_clustering")

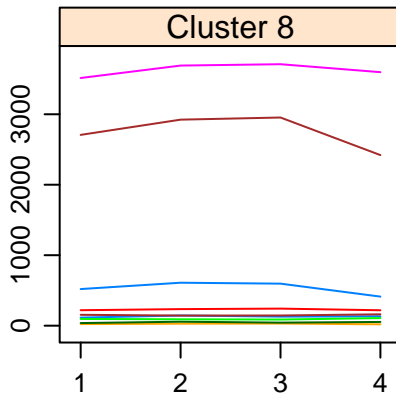
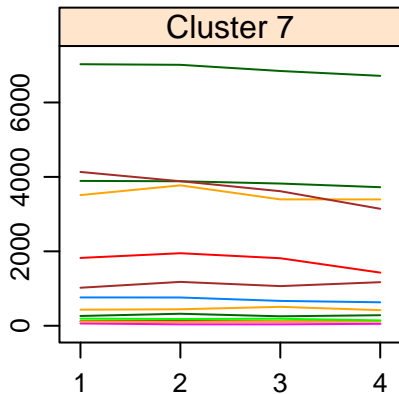
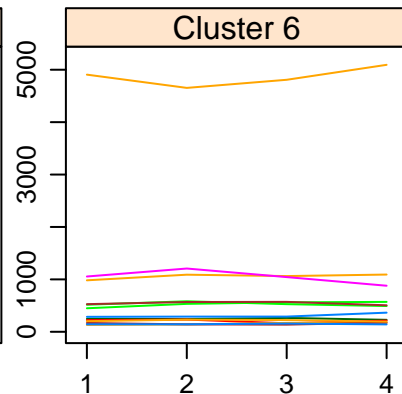
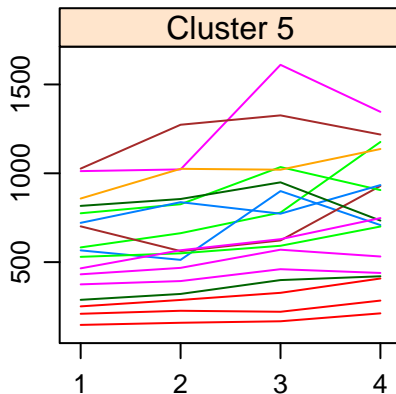
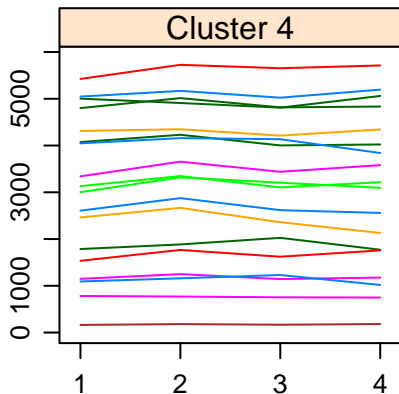
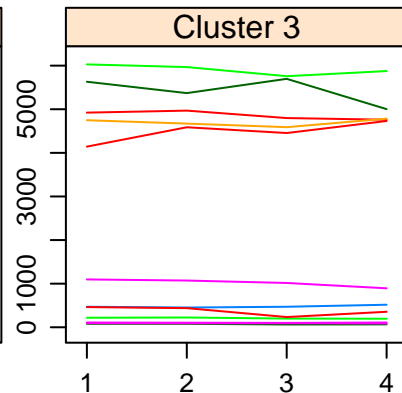
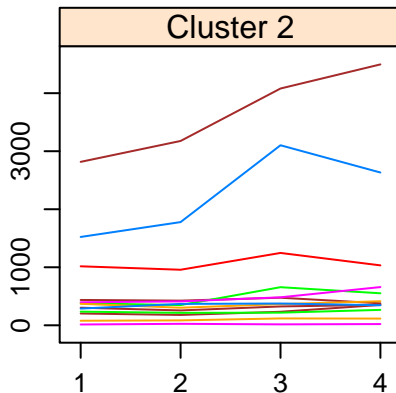
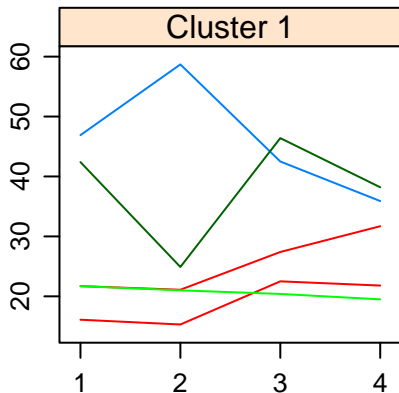
Subject 5



help("unsupervised_clustering")

Subject 6

Gene Expression



help("unsupervised_clustering")

