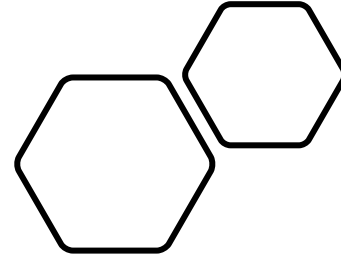
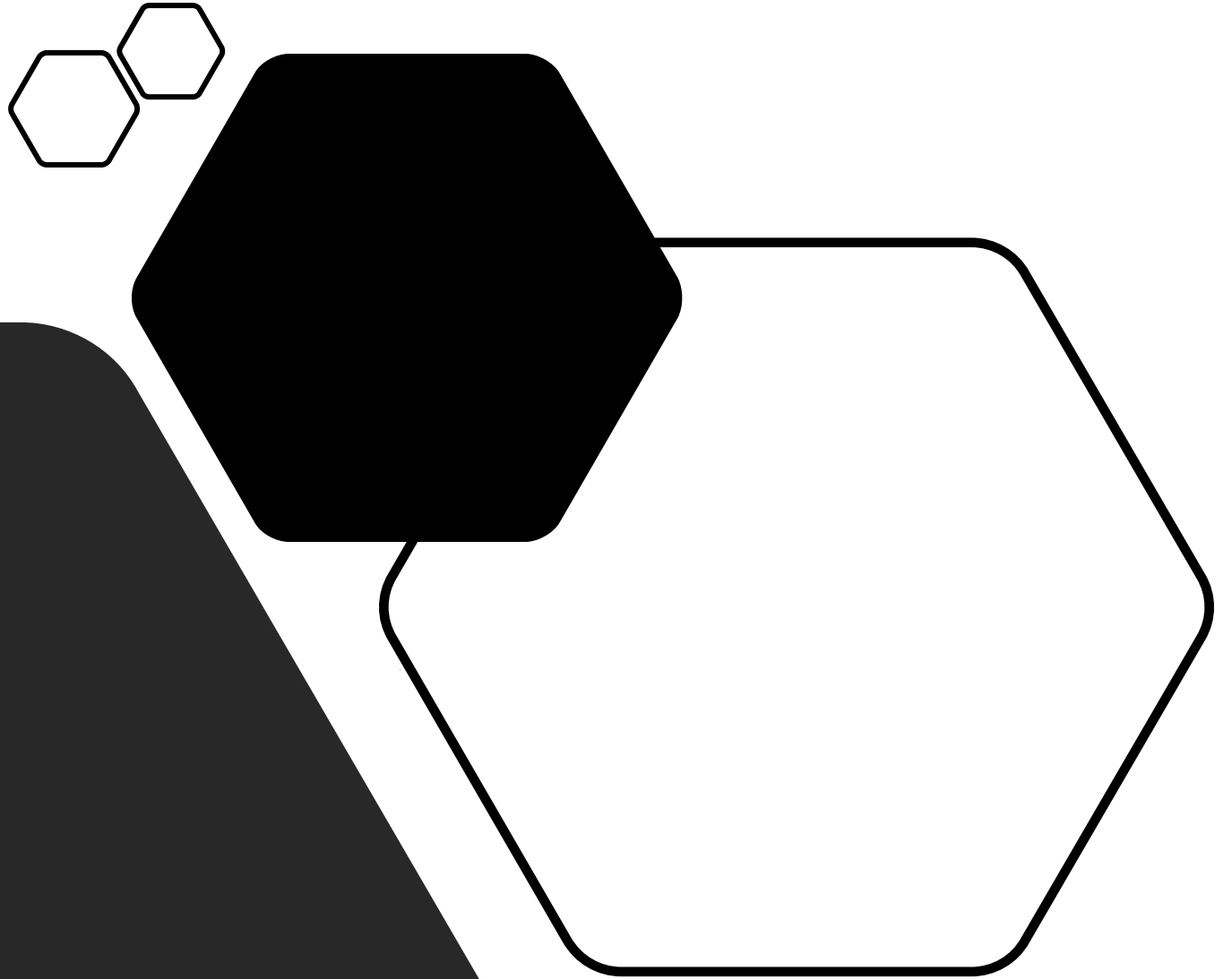


# Bioinformatic Analysis CD151, EWI2, TSPAN8



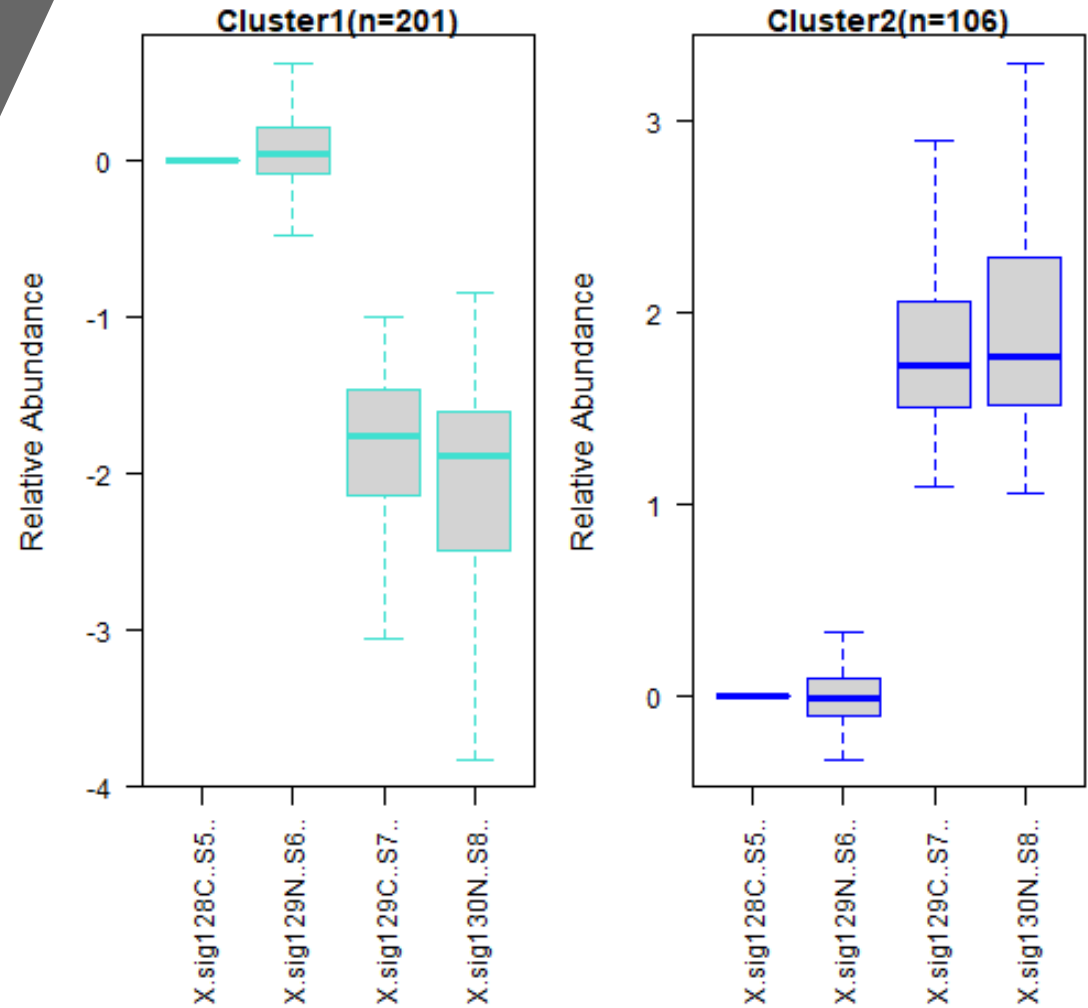
David R Vanderwall

# EWI2 Knockout

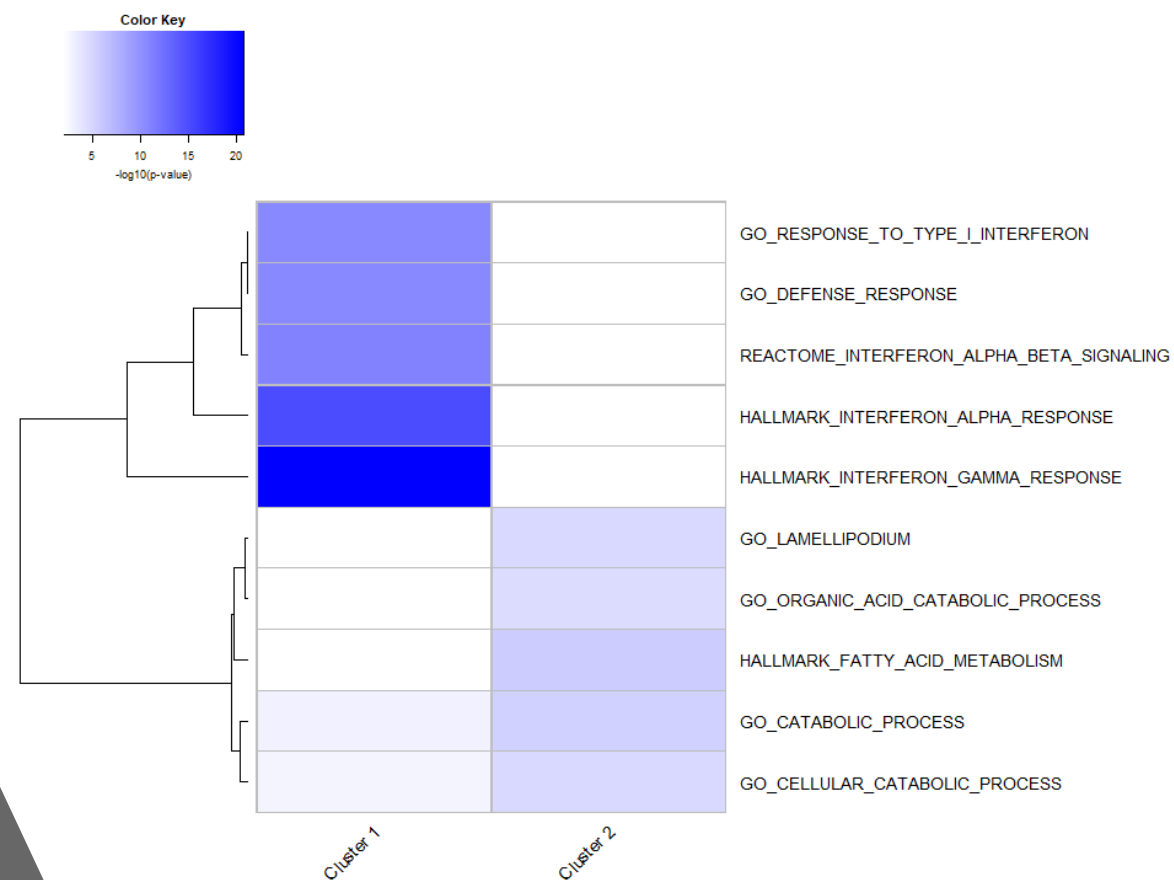


# WGCNA Co-Expression Clustering

Upregulated and Downregulated Clusters identified



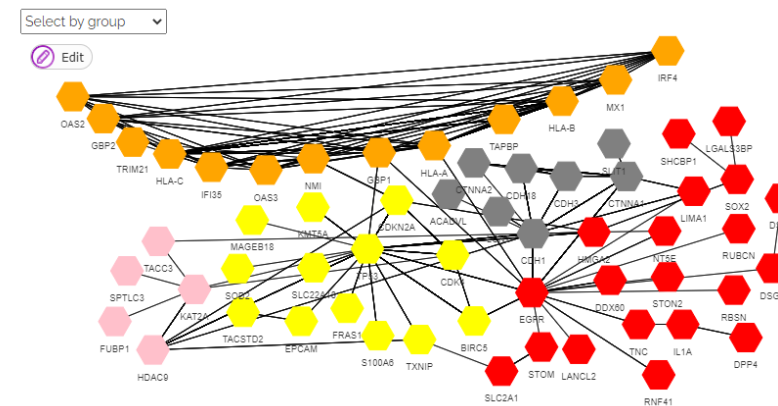
# Cluster Pathway Analysis



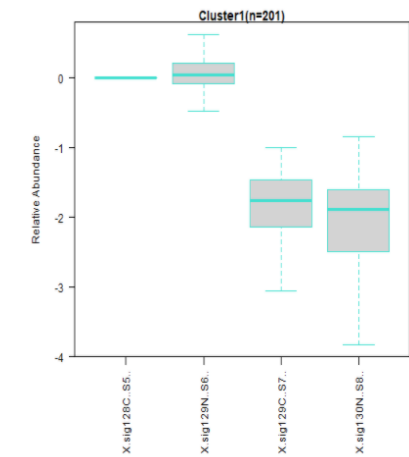
# PPI

## Modularization of Downregulated Cluster

Co-Expression Cluster 1: 11 Modules Found, 73  
Constituent Genes

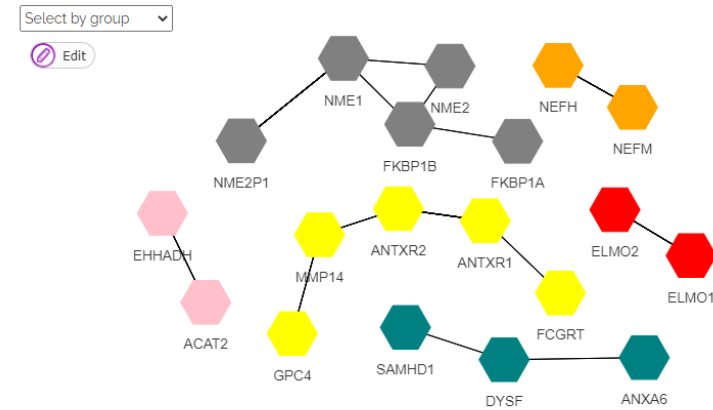


Cluster 1 Boxplot Trend



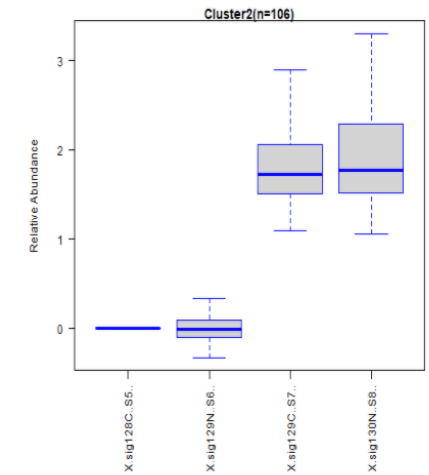
# PPI Modularization of Upregulated Cluster

Co-Expression Cluster 2: 6 Modules Found, 19 Constituent Genes



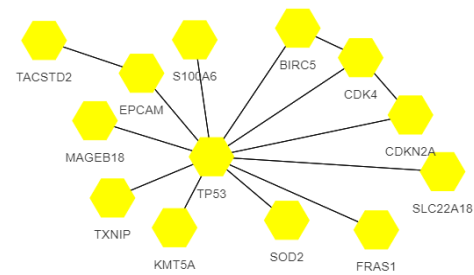
Select a Network Layout Format

Cluster 2 Boxplot Trend

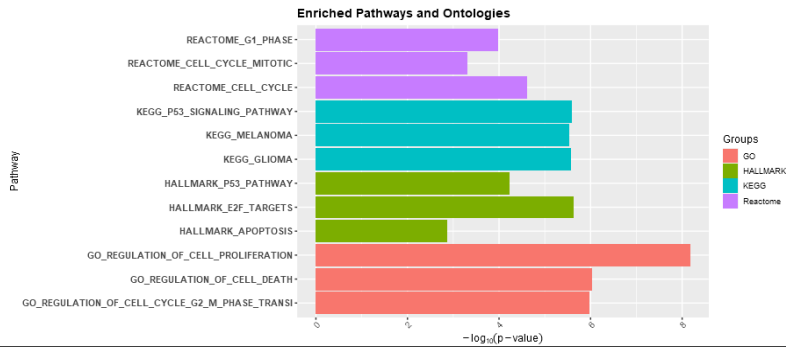


Cluster1\_Module2: 13 Constituent Genes

Edit

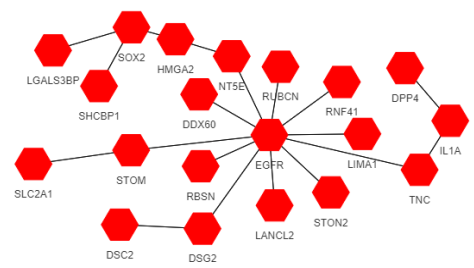


Cluster1\_Module2 Pathway Barplot

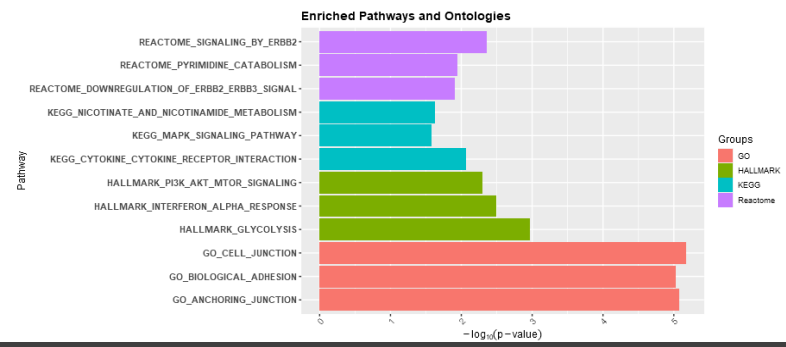


Cluster1\_Module1: 20 Constituent Genes

Edit



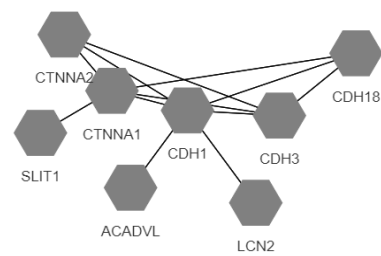
Cluster1\_Module1 Pathway Barplot



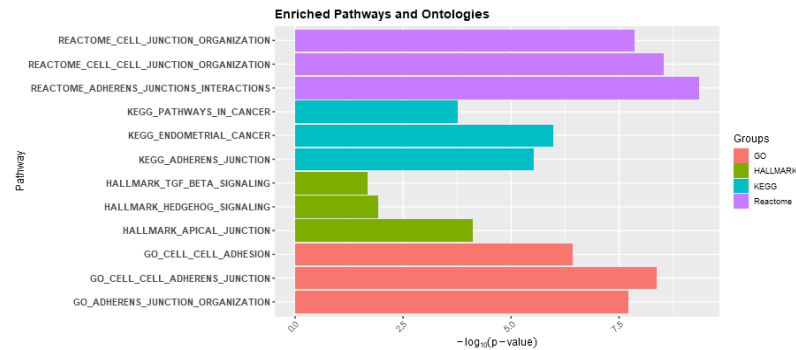
# Pathway Analysis Part I

Cluster1\_Module4: 8 Constituent Genes

Edit

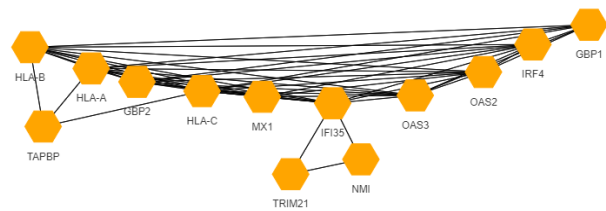


Cluster1\_Module4 Pathway Barplot

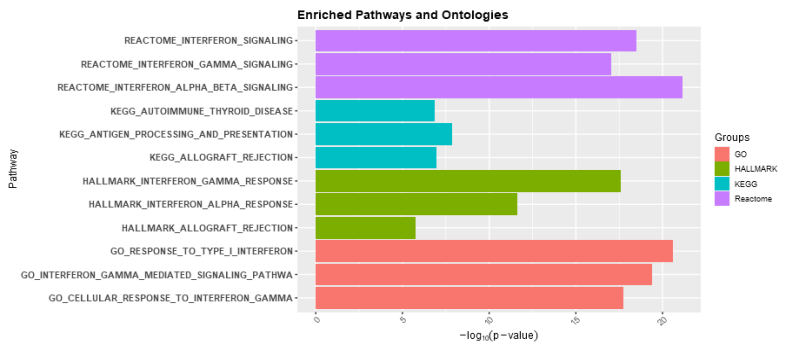


Cluster1\_Module3: 13 Constituent Genes

Edit



Cluster1\_Module3 Pathway Barplot

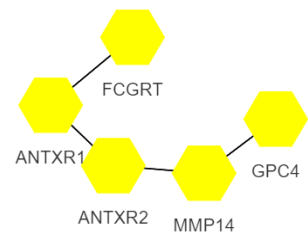


# Pathway Analysis Part II

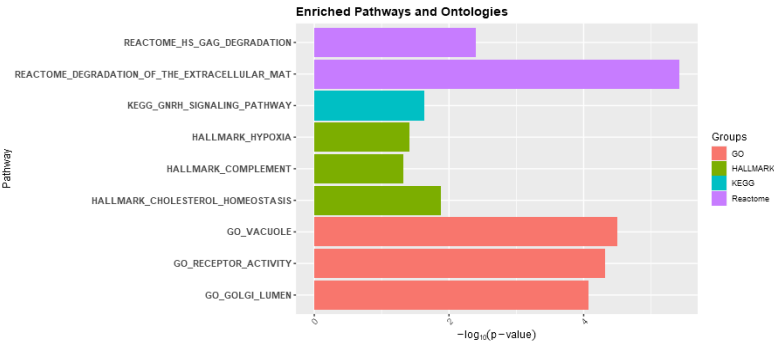


Cluster2\_Module2: 5 Constituent Genes

Edit

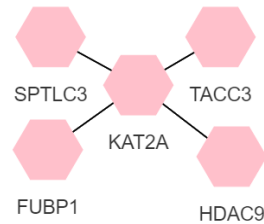


Cluster2\_Module2 Pathway Barplot

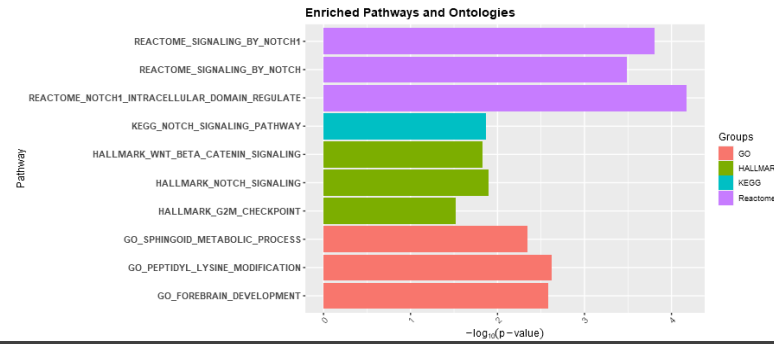


Cluster1\_Module5: 5 Constituent Genes

Edit

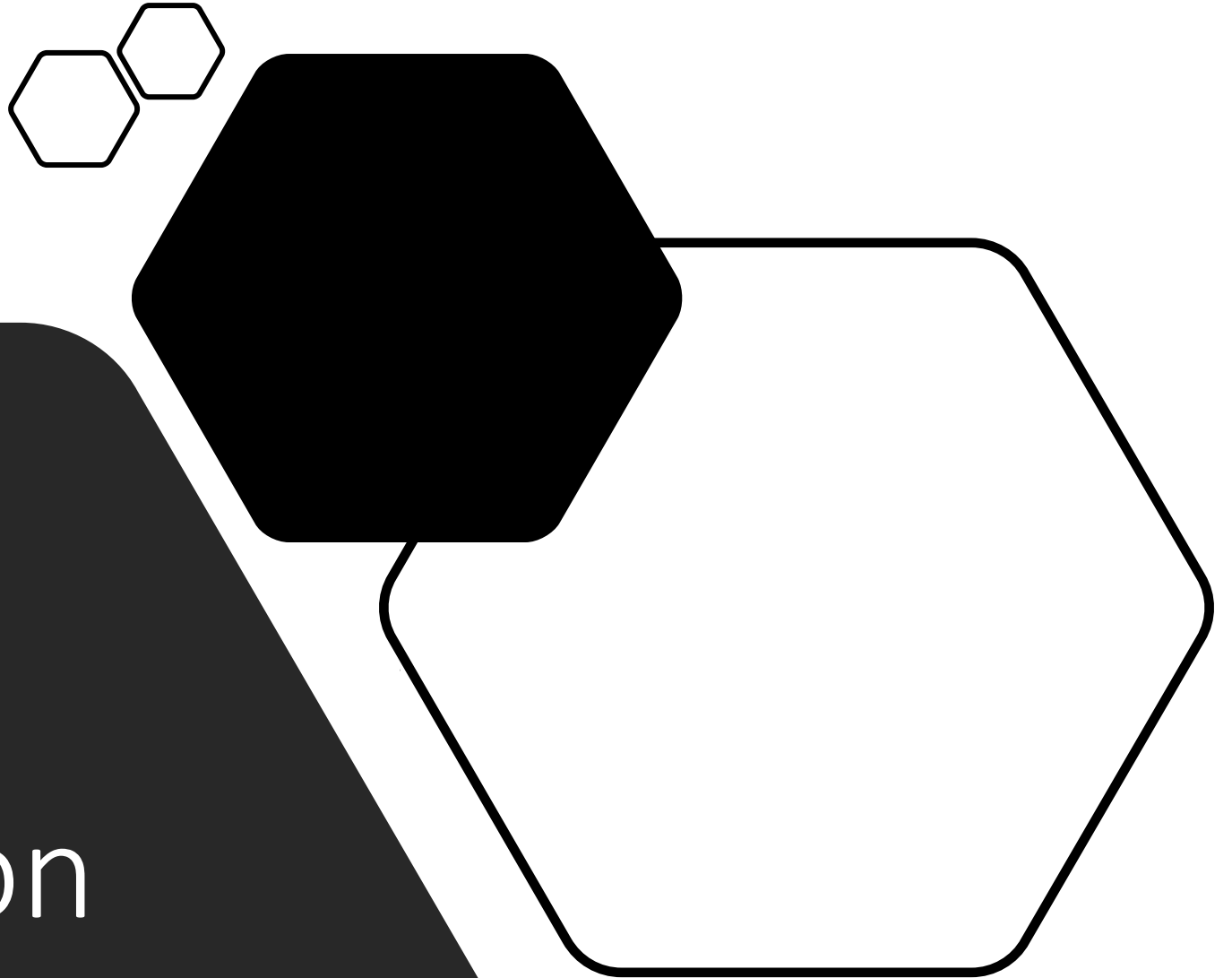


Cluster1\_Module5 Pathway Barplot



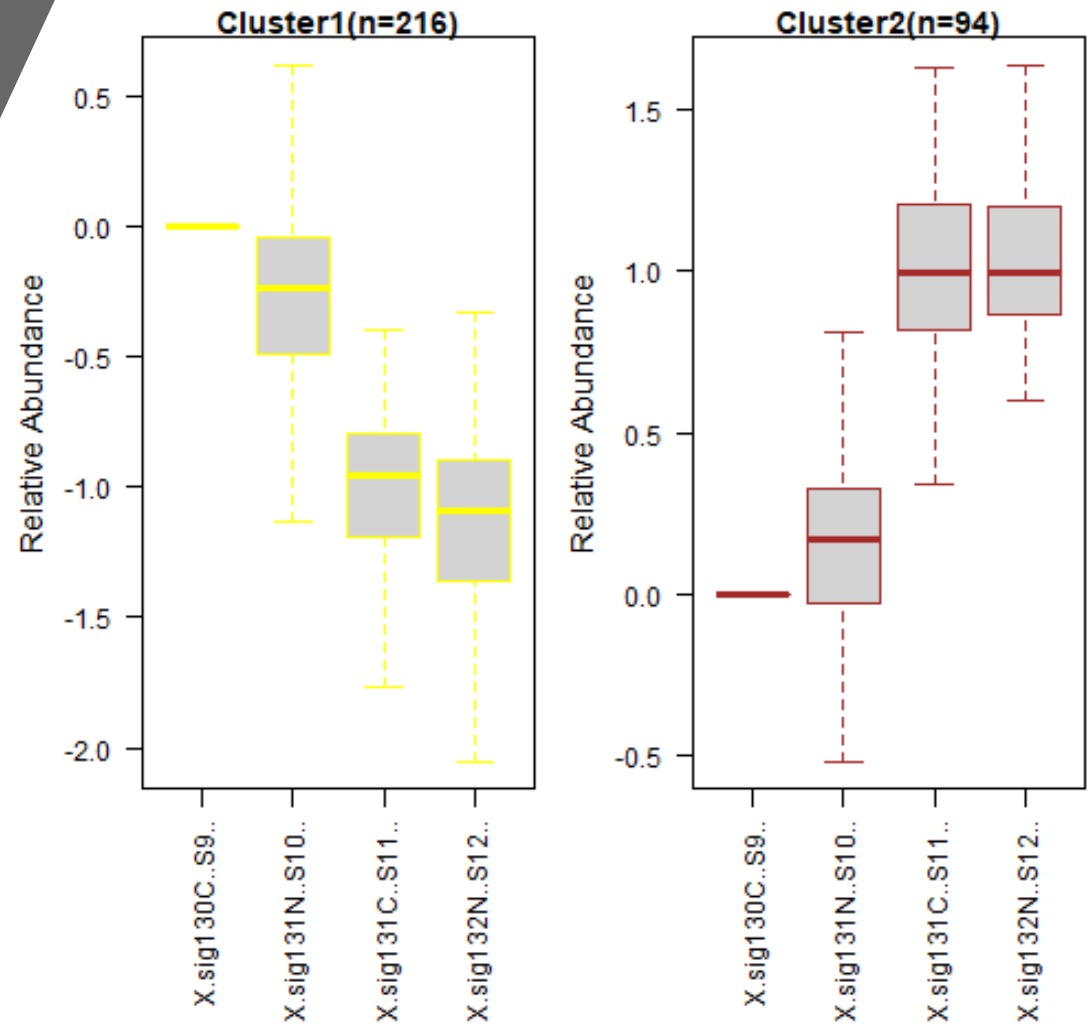
Pathway Analysis Part III

EWI2  
Overexpression

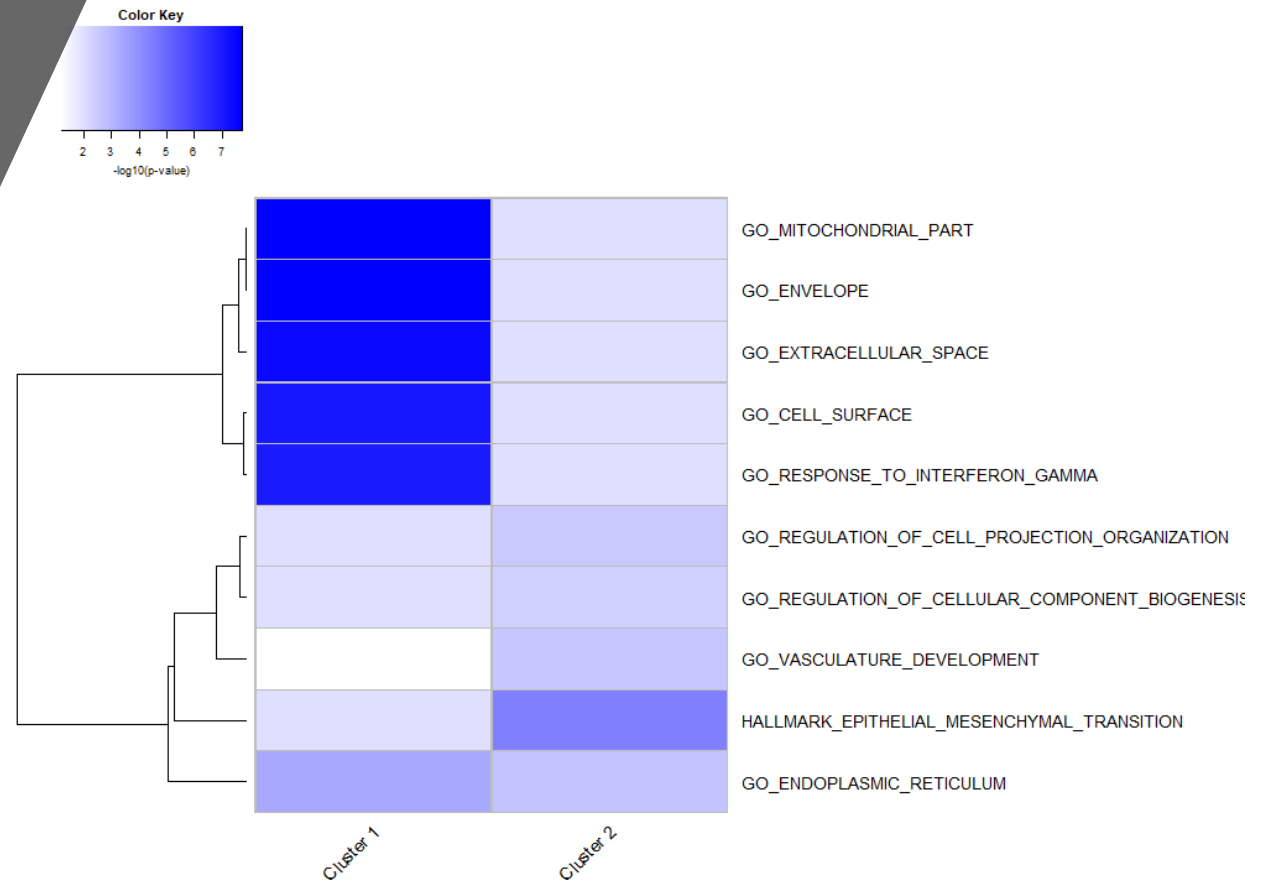


# WGCNA Co-Expression Clustering

Upregulated and Downregulated Clusters identified

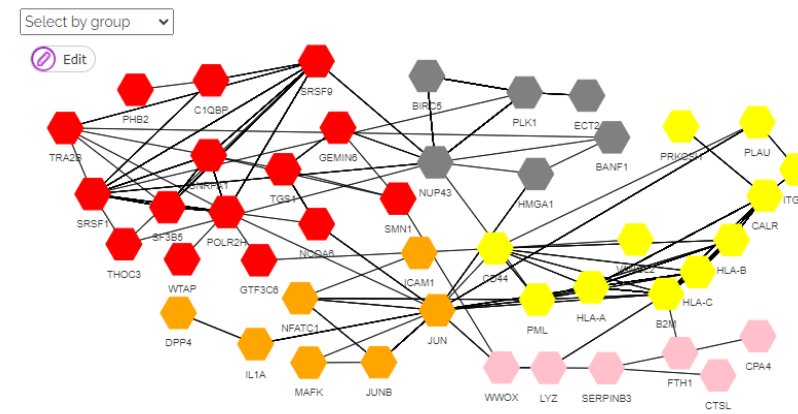


# Cluster Pathway Analysis

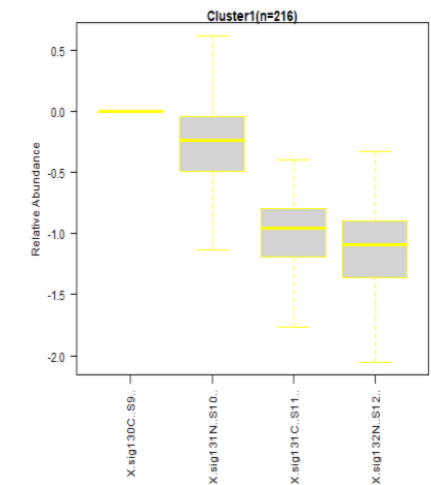


# PPI Modularization of Downregulated Cluster

Co-Expression Cluster 1: 13 Modules Found, 68  
Constituent Genes



Cluster 1 Boxplot Trend

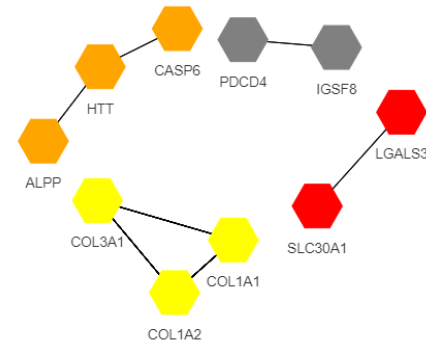


# PPI Modularization of Upregulated Cluster

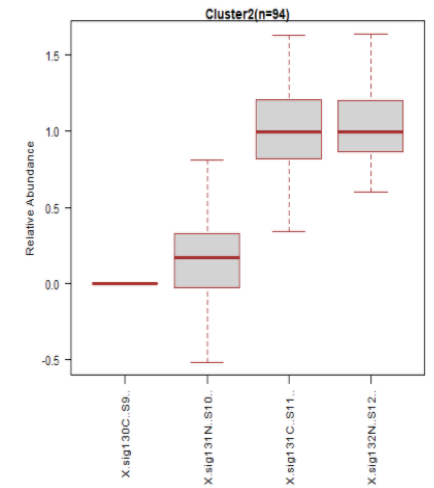
Co-Expression Cluster 2: 4 Modules Found, 10 Constituent Genes

Select by group

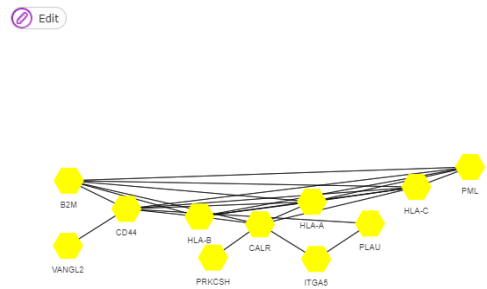
Edit



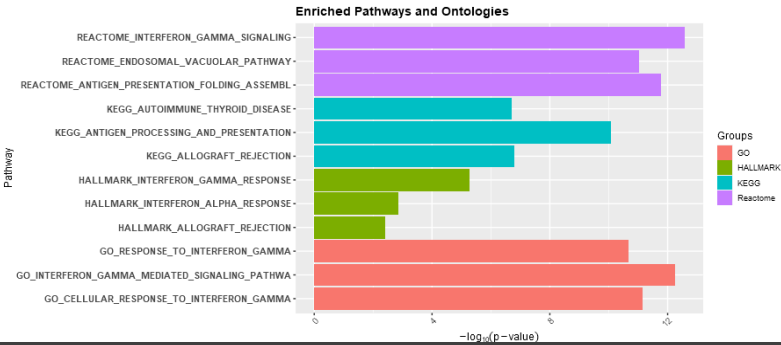
Cluster 2 Boxplot Trend



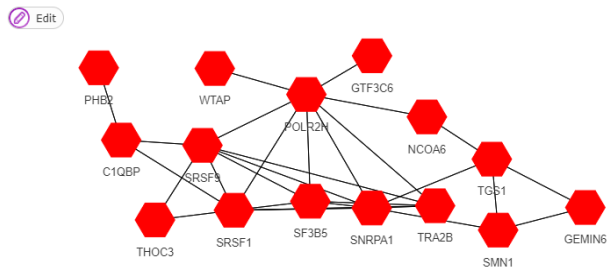
Cluster1\_Module2: 11 Constituent Genes



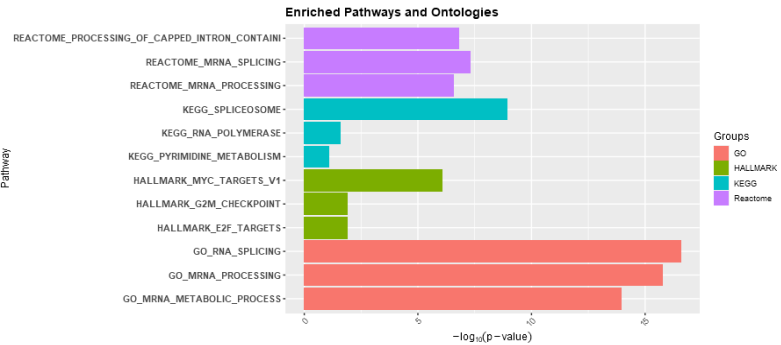
Cluster1\_Module2 Pathway Barplot



Cluster1\_Module1: 15 Constituent Genes



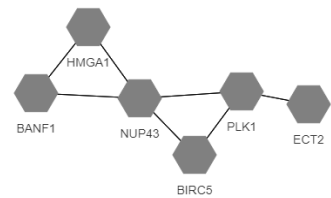
Cluster1\_Module1 Pathway Barplot



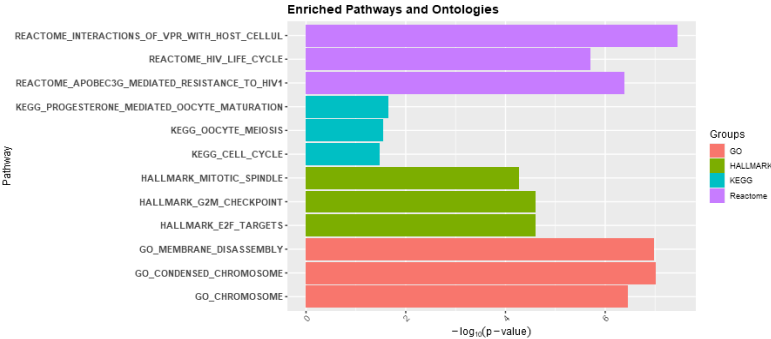
# Pathway Analysis Part I

Cluster1\_Module4: 6 Constituent Genes

Edit

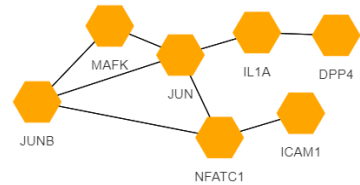


Cluster1\_Module4 Pathway Barplot

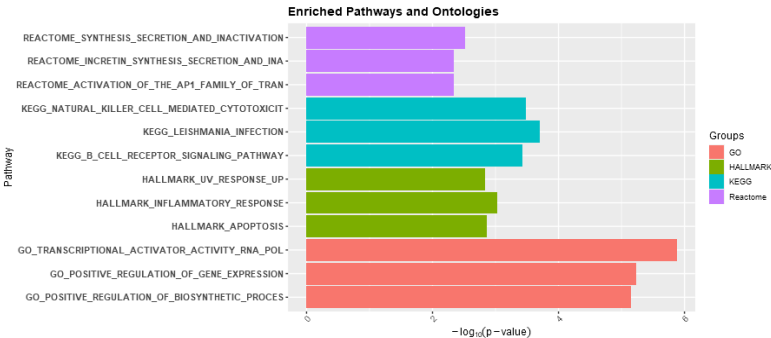


Cluster1\_Module3: 7 Constituent Genes

Edit



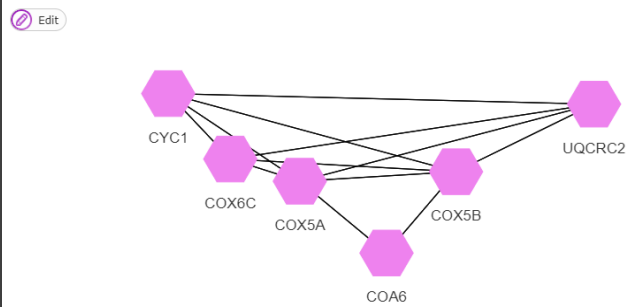
Cluster1\_Module3 Pathway Barplot



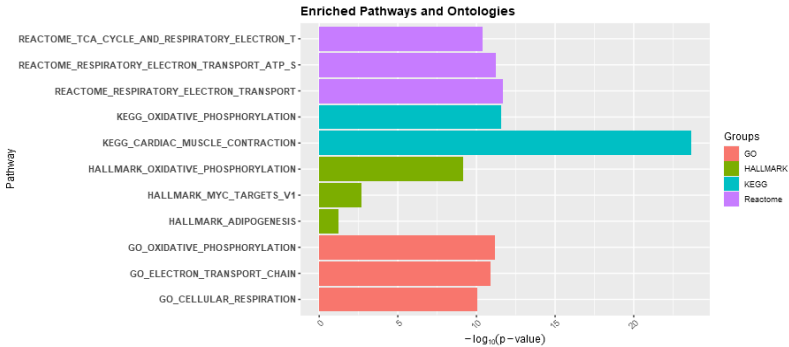
Pathway Analysis Part II



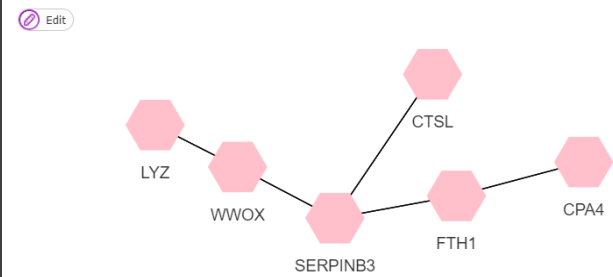
Cluster1\_Module7: 6 Constituent Genes



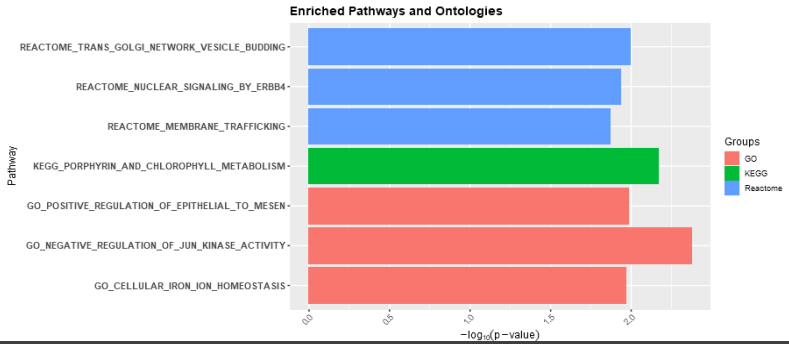
Cluster1\_Module7 Pathway Barplot



Cluster1\_Module5: 6 Constituent Genes

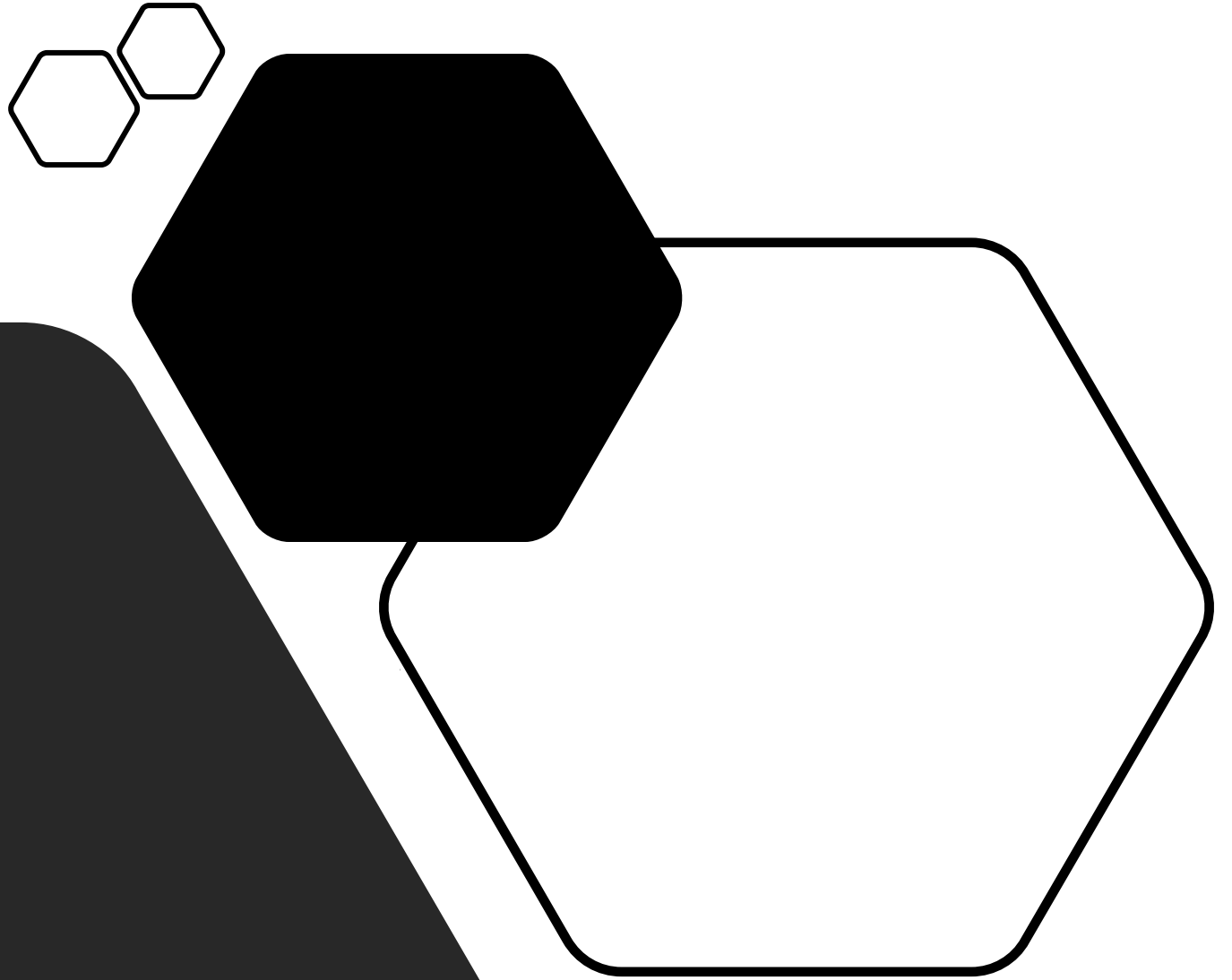


Cluster1\_Module5 Pathway Barplot



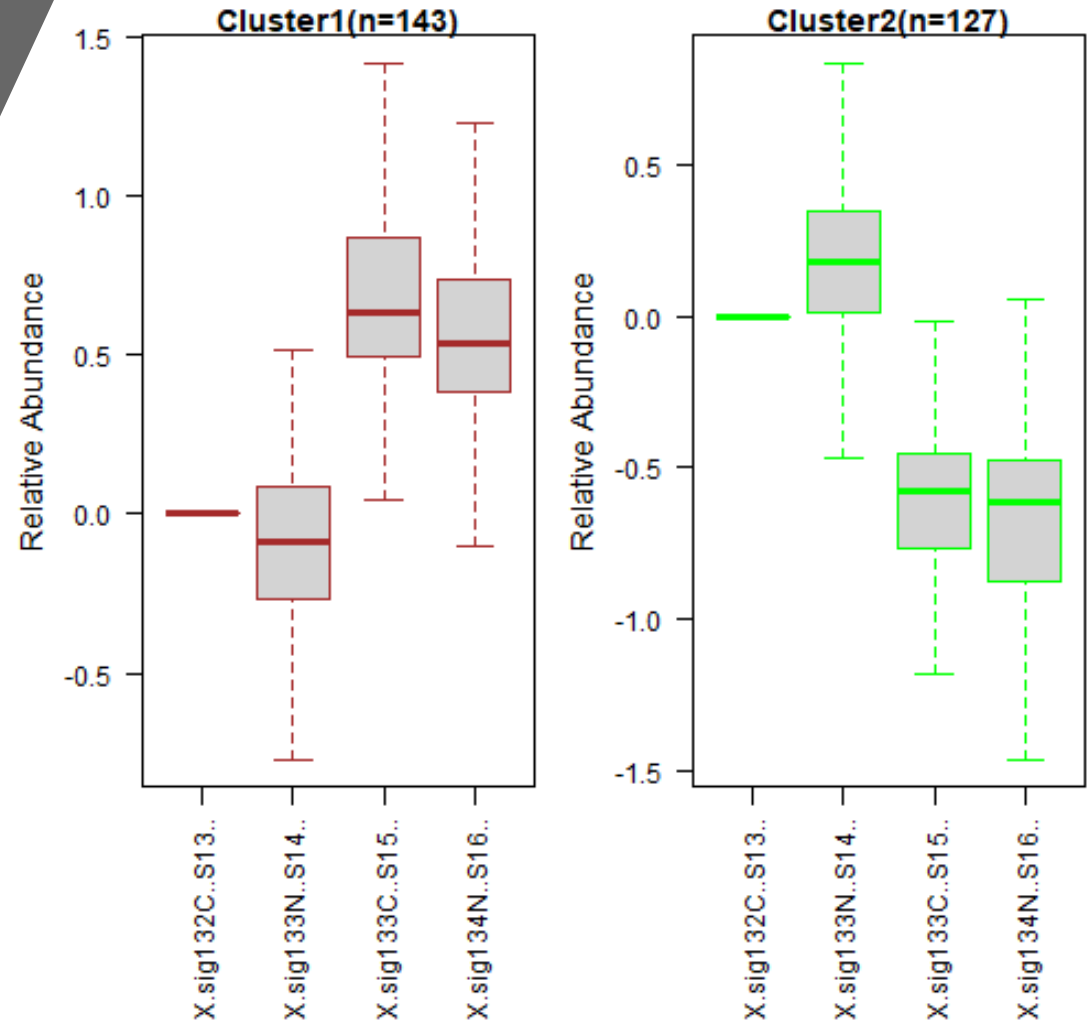
# Pathway Analysis Part III

# TSPAN8 Knockdown

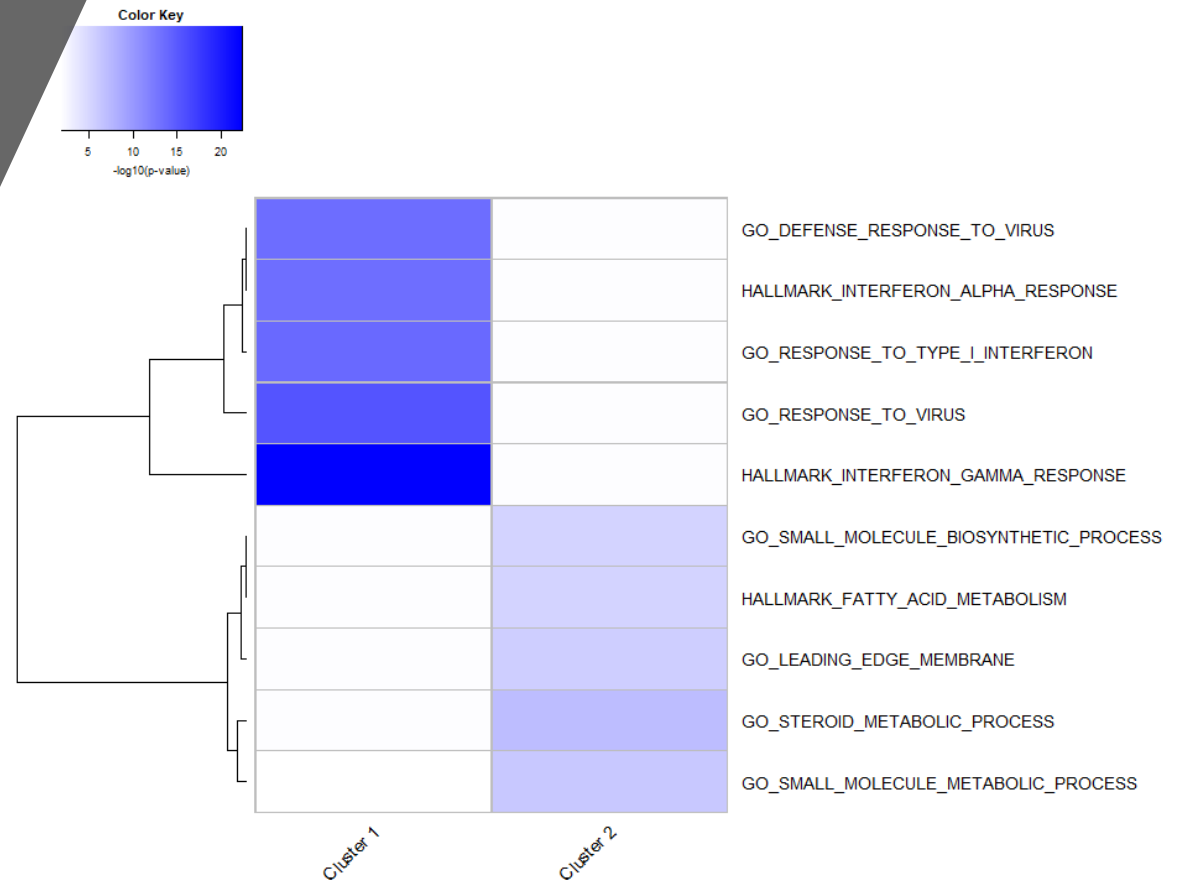


# WGCNA Co-Expression Clustering

Upregulated and Downregulated Clusters identified

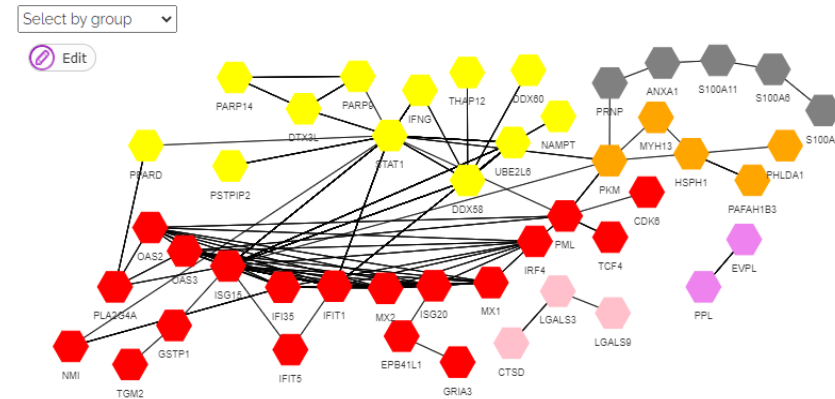


# Cluster Pathway Analysis

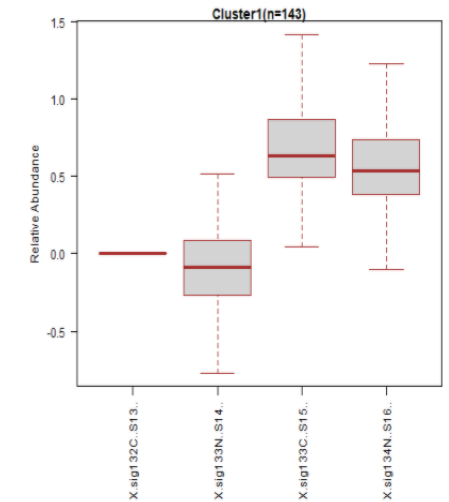


# PPI Modularization of Upregulated Cluster

Co-Expression Cluster 1: 7 Modules Found, 48 Constituent Genes

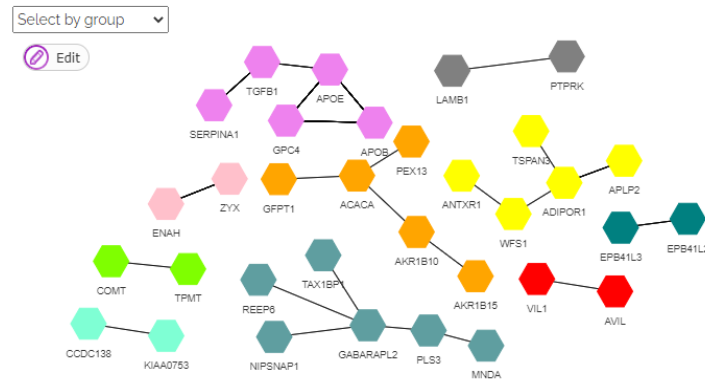


Cluster 1 Boxplot Trend

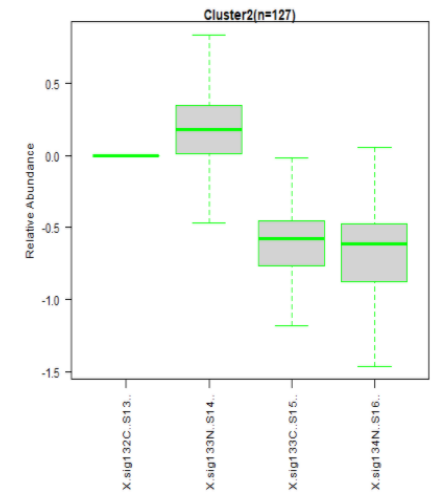


# PPI Modularization of Downregulated Cluster

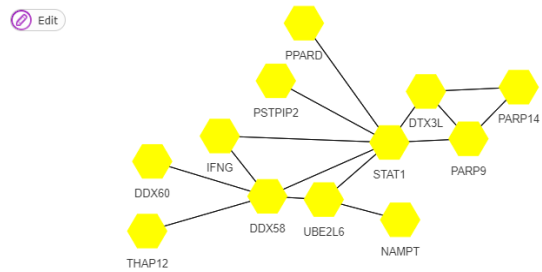
Co-Expression Cluster 2: 10 Modules Found, 33  
Constituent Genes



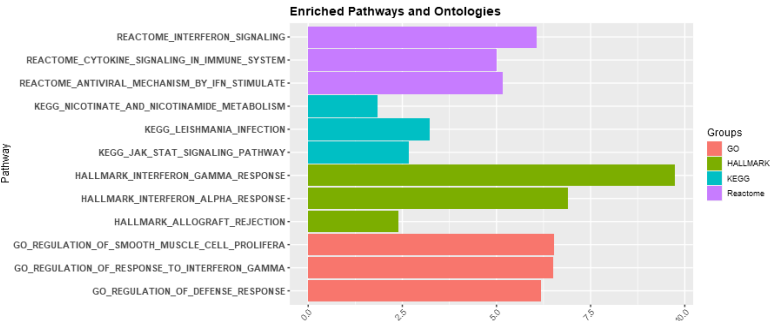
Cluster 2 Boxplot Trend



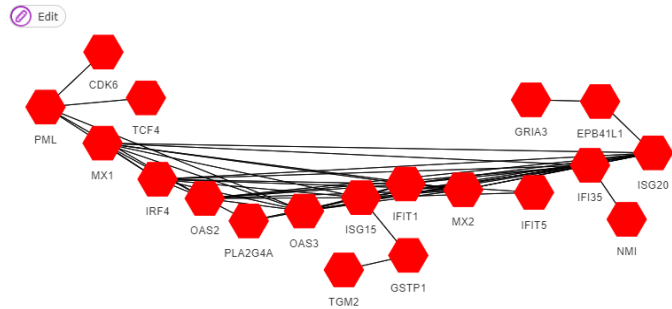
Cluster1\_Module2: 12 Constituent Genes



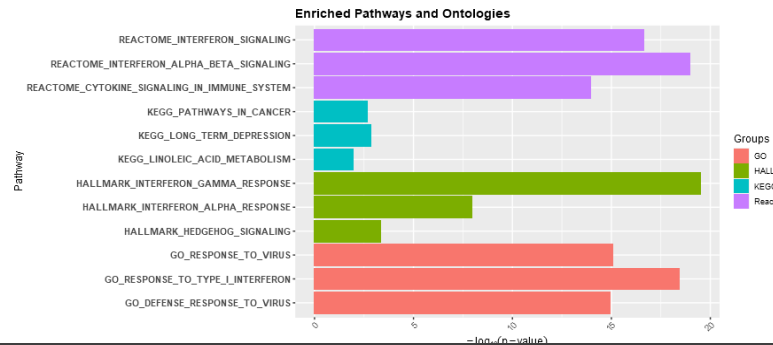
Cluster1\_Module2 Pathway Barplot



Cluster1\_Module1: 19 Constituent Genes



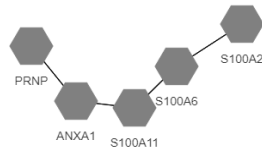
Cluster1\_Module1 Pathway Barplot



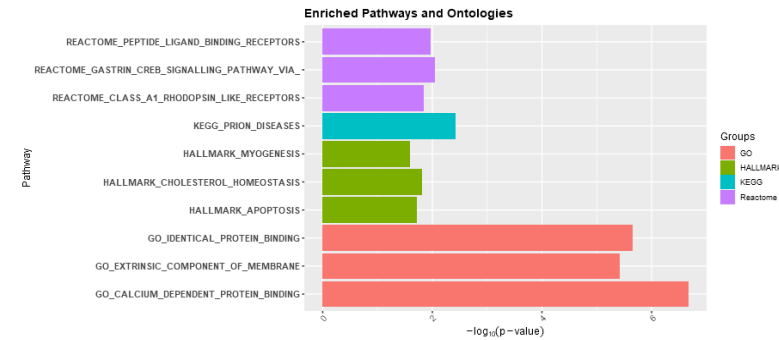
# Pathway Analysis Part I

### Cluster1\_Module4: 5 Constituent Genes

Edit

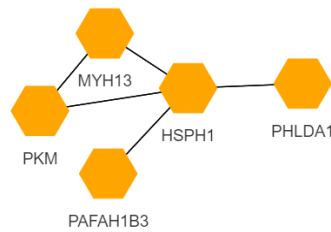


### Cluster1\_Module4 Pathway Barplot

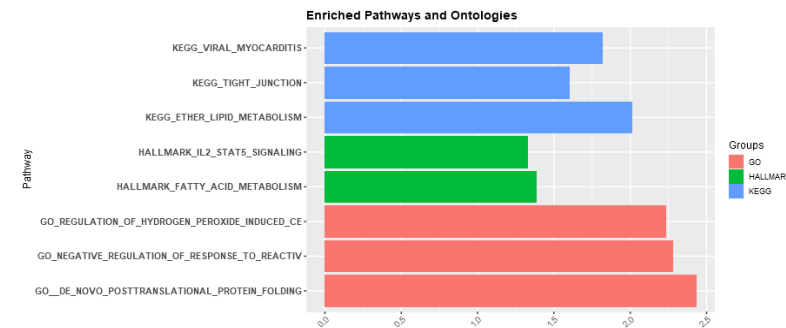


### Cluster1\_Module3: 5 Constituent Genes

Edit



### Cluster1\_Module3 Pathway Barplot

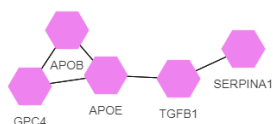


# Pathway Analysis Part II

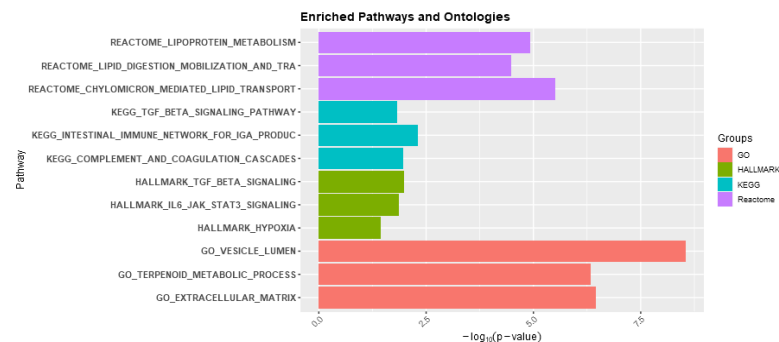


### Cluster2\_Module7: 5 Constituent Genes

Edit

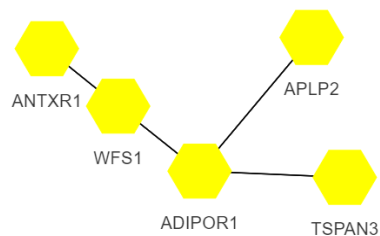


### Cluster2\_Module7 Pathway Barplot

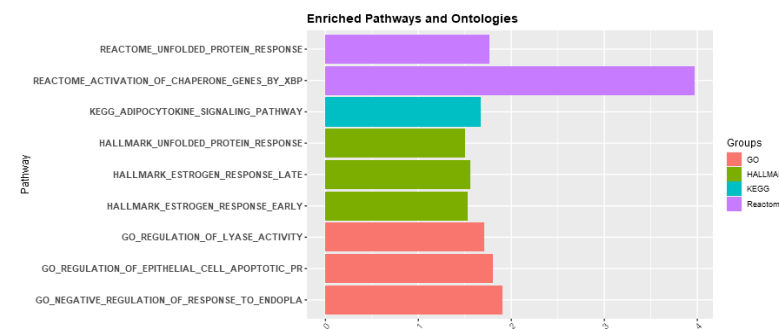


### Cluster2\_Module2: 5 Constituent Genes

Edit

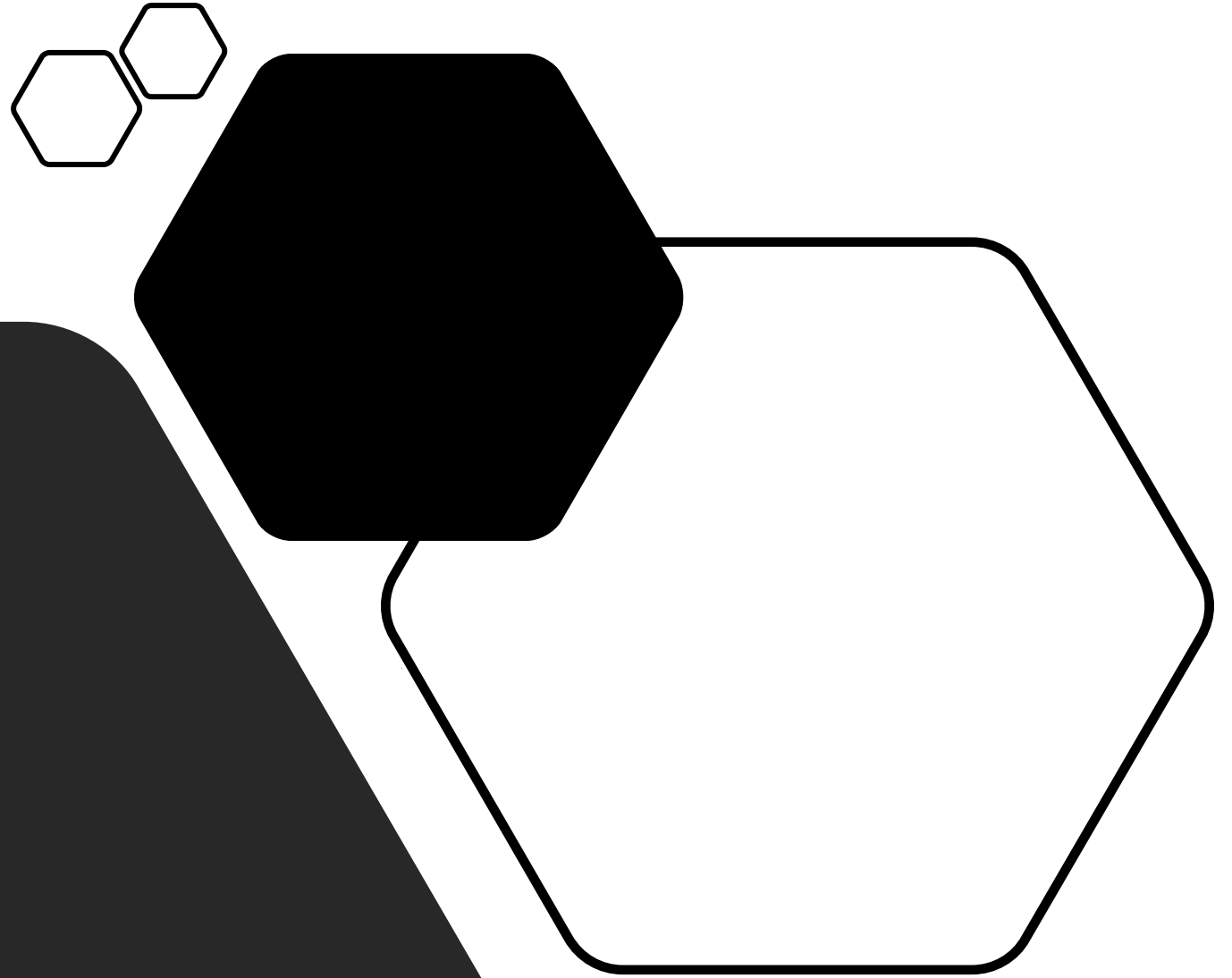


### Cluster2\_Module2 Pathway Barplot



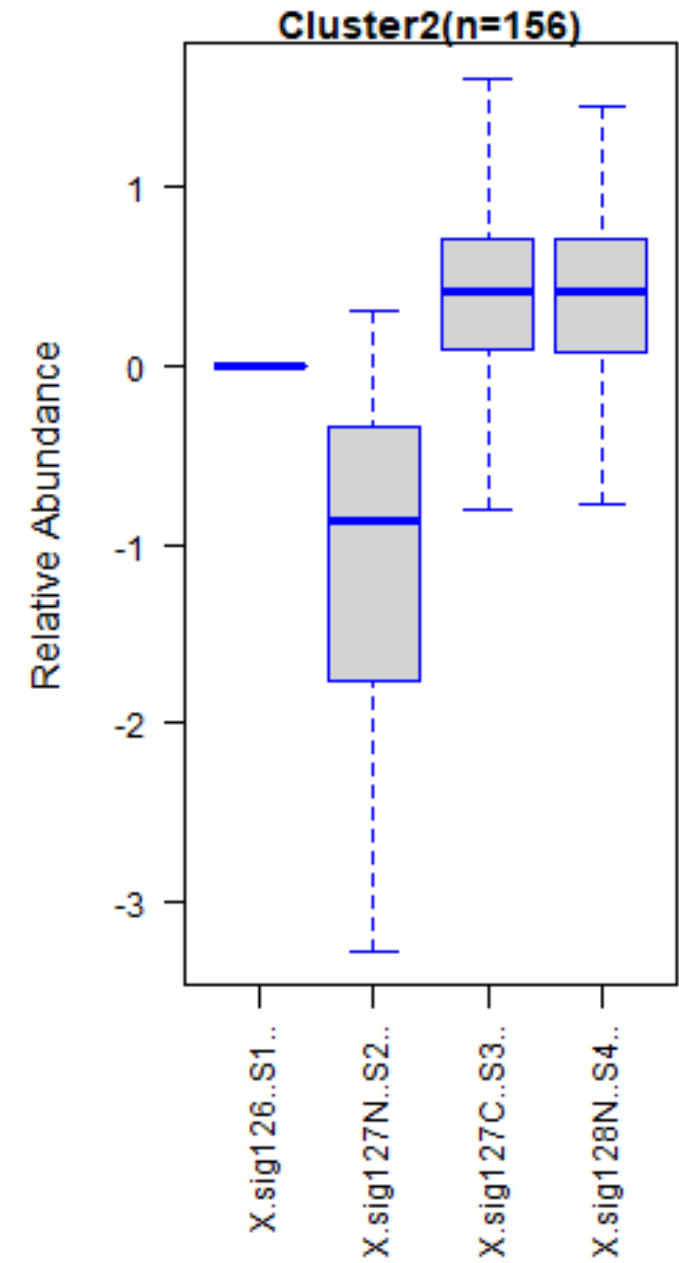
# Pathway Analysis Part III

# CD151 Knockdown

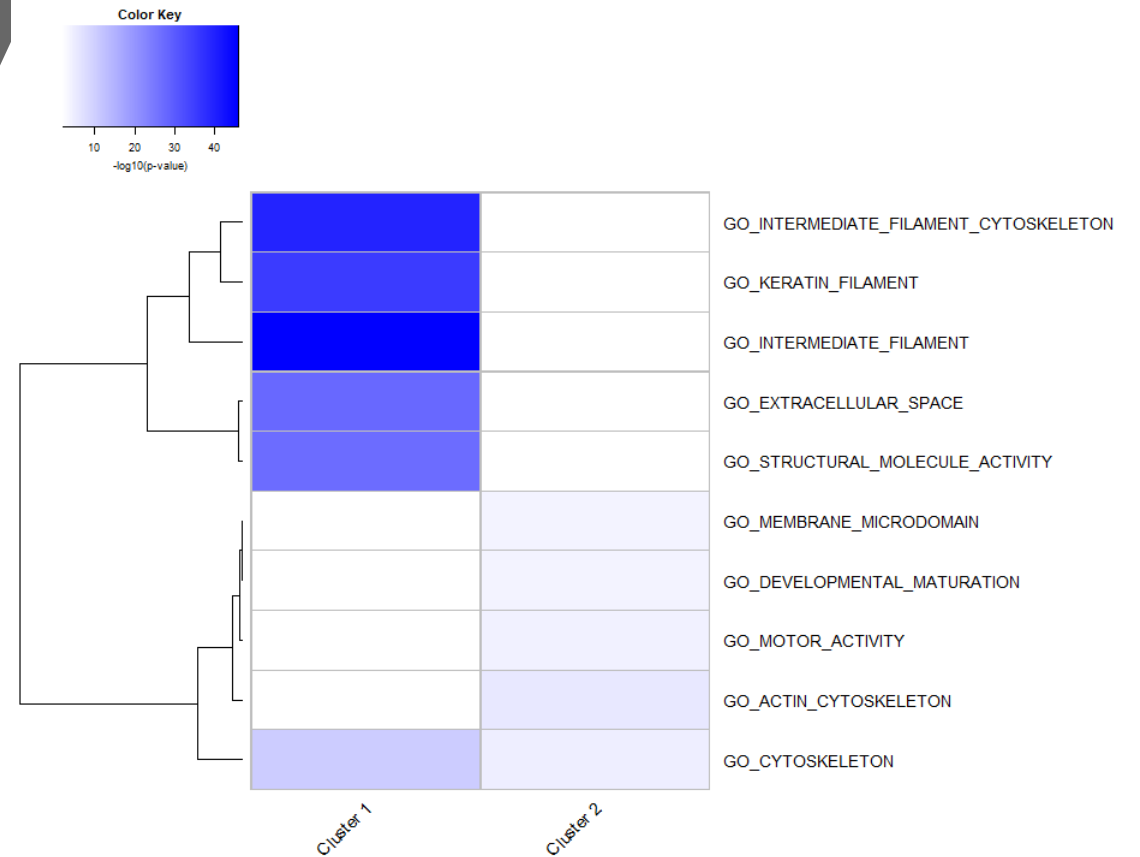


# WGCNA Co-Expression Clustering

Upregulated and  
Downregulated Clusters  
identified

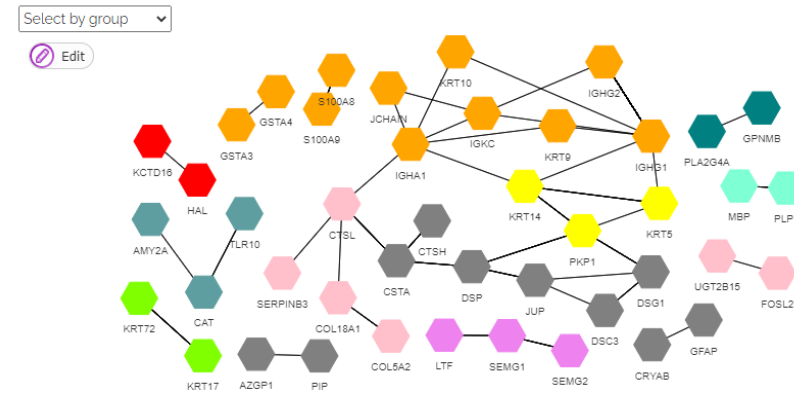


# Cluster Pathway Analysis

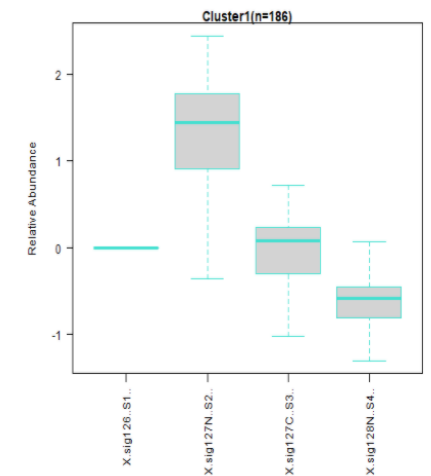


# PPI Modularization of Downregulated Cluster

Co-Expression Cluster 1: 15 Modules Found, 44  
Constituent Genes

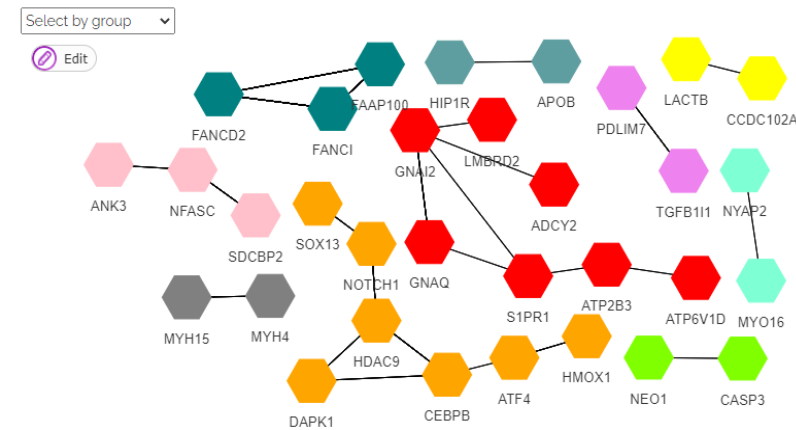


Cluster 1 Boxplot Trend

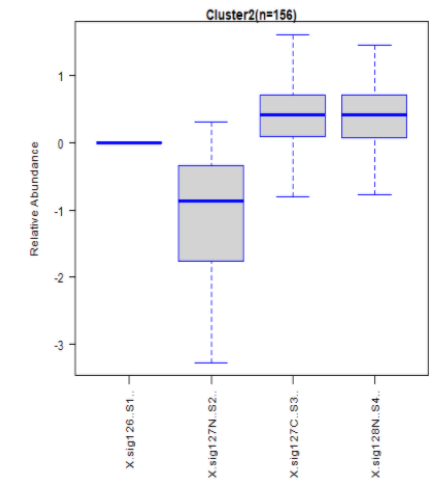


# PPI Modularization of Upregulated Cluster

Co-Expression Cluster 2: 10 Modules Found, 32  
Constituent Genes

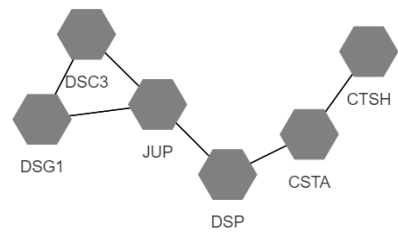


Cluster 2 Boxplot Trend

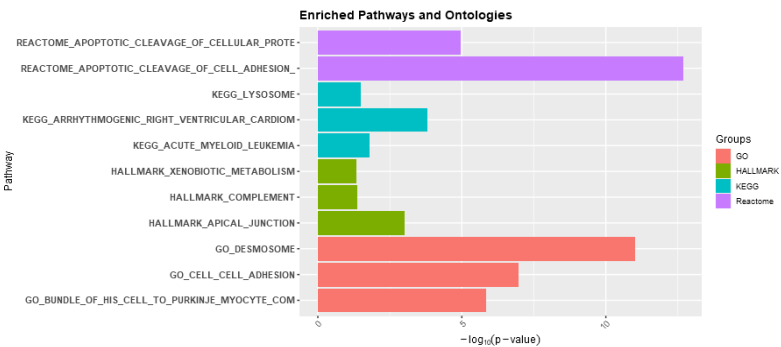


Cluster1\_Module5: 6 Constituent Genes

Edit

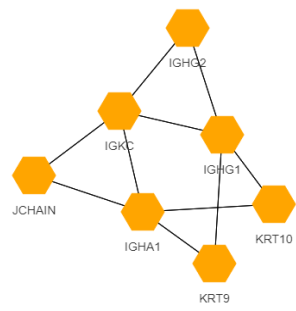


Cluster1\_Module5 Pathway Barplot

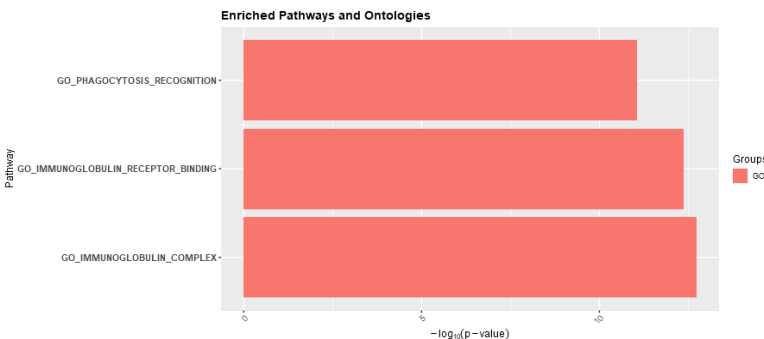


Cluster1\_Module4: 7 Constituent Genes

Edit

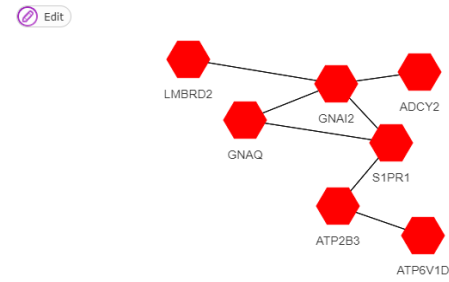


Cluster1\_Module4 Pathway Barplot

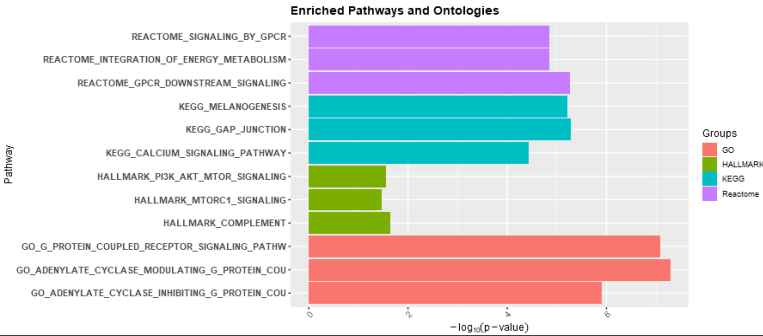


Pathway Analysis Part I

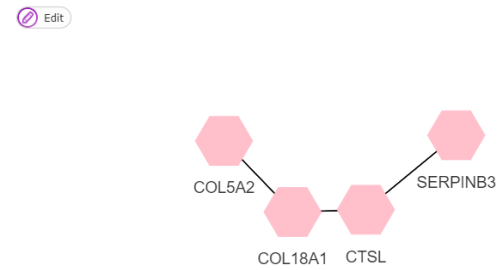
Cluster2\_Module1: 7 Constituent Genes



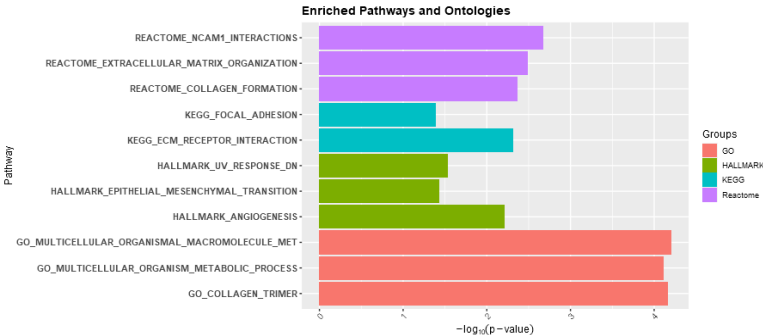
Cluster2\_Module1 Pathway Barplot



Cluster1\_Module6: 4 Constituent Genes



Cluster1\_Module6 Pathway Barplot

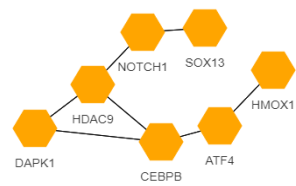


# Pathway Analysis Part II

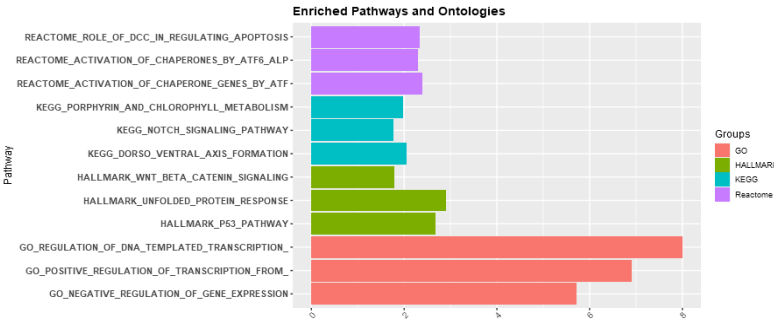


Cluster2\_Module3: 7 Constituent Genes

Edit



Cluster2\_Module3 Pathway Barplot

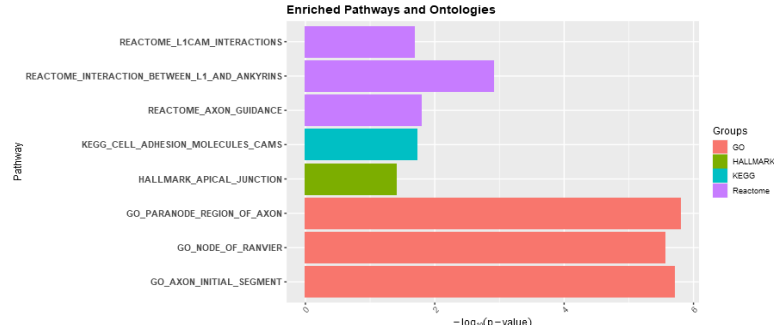


Cluster2\_Module5: 3 Constituent Genes

Edit



Cluster2\_Module5 Pathway Barplot



Pathway Analysis Part III