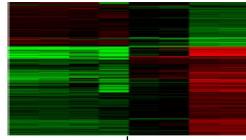


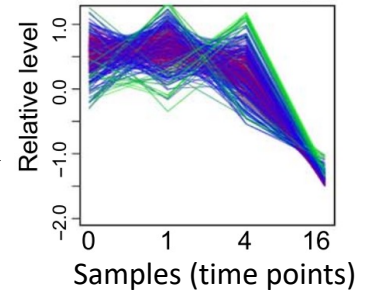
Quantification matrix of
variable or DE proteins



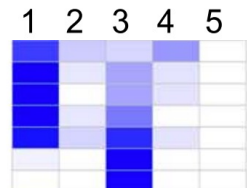
Outputs

Co-expression clusters

Cluster 1
(n = 411 proteins)



Co-expression clusters



Significantly
enriched
pathways

Annotated protein modules

Cluster 1: Module 5 (proteasome)



JUMPn modules

Co-expression
clustering analysis
(WGCNA)

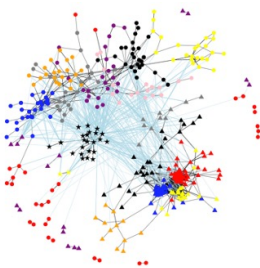
Pathway
enrichment analysis

Protein-protein
interaction network
analysis

Public
databases



PPI network



Publication tables
in Microsoft Excel format