

Supplemental.1

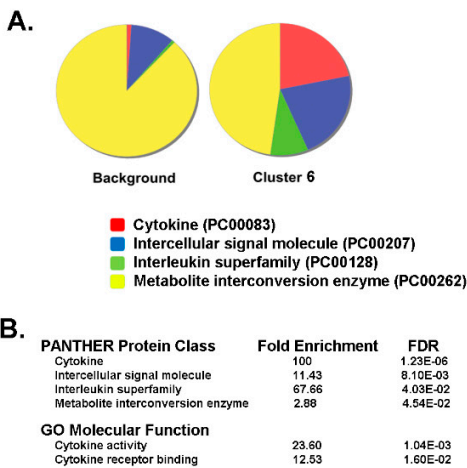


Figure S1. Enrichment analysis of the DEGs in Cluster 6. **(A)** Pie charts of protein class variation after a GO enrichment analysis showing the behavior of background genes and genes clustered in 6. **(B)** Fold enrichment values for protein class and molecular functions with a FDR ≤ 0.05 as a cut off, and using as background all genes detected.

Supplemental.2

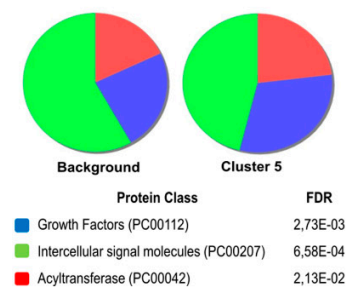


Figure S2. Gene ontology (GO) enrichment analysis results of genes from cluster 5. Pie chart of protein class variation with a false discovery rate (FDR) ≤ 0.05 as a cut off, and using all genes detected as background.