Overrepresentation: GO term (Biological process) for all significant cluster 12 marker genes ribonucleoprotein complex biogenesis mRNA processing ncRNA processing RNA splicingproteasomal protein catabolic process regulation of DNA metabolic process histone modificationregulation of translation establishment of protein localization to organelle mitotic cell cycle phase transition proteasome-mediated ubiquitin-dependent proteincatabolic process ribosome biogenesis non-membrane-bounded Count organelle assembly 150 RNA splicing, via 200 transesterification reactions 250 RNA splicing, via transesterification 300 reactions with bulged adenosine as nucleophile p.adjust mRNA splicing, via spliceosome 5.376201e-29 4.032151e-29 chromosome segregation -2.688101e-29 1.344050e-29 rRNA metabolic process 1.319494e-85 regulation of mRNA metabolic process nucleocytoplasmic transportnuclear transport-DNA replication mitotic nuclear division -RNA catabolic processrRNA processingregulation of chromosome organization mRNA catabolic process sister chromatid segregation RNA localization cytoplasmic translation -0.03 0.04 0.05 0.06 GeneRatio