Overrepresentation: GO term (Biological process) for top 100 significant cluster 12 marker genes cytoplasmic translation ribonucleoprotein complex biogenesis ribosome biogenesis ncRNA processing non-membrane-bounded organelle assembly rRNA processing rRNA metabolic process ribosome assembly ribonucleoprotein complex assembly ribonucleoprotein complex subunit organization regulation of translation ribosomal large subunit biogenesis ribosomal small subunit Count biogenesis 10 positive regulation of translation positive regulation of cellular amide metabolic process p.adjust cell killing 0.003 regulation of protein 0.002 ubiquitination 0.001 ribosomal small subunit assembly ribosomal large subunit assembly maturation of SSU-rRNA regulation of signal transduction by p53 class mediator translational initiation maturation of SSU-rRNA from tricistronic rRNA transcript (SSU-rRNA, 5.8S rRNA, LSU-rRNA) negative regulation of protein ubiquitination negative regulation of protein modification by small protein conjugation or positive regulation of signal transduction by p53 class mediator regulation of intrinsic apoptotic signaling pathway in response to DNA damage negative regulation of ubiquitin protein ligase antigen processing and presentation of endogenous peptide antigen via MHC antigen processing and presentation of endogenous peptide antigen 0.2 0.4 0.6 GeneRatio