Number of genes
20
0.50
0.25
adaptive immune response based on somatic recombination of immune receptors built from immunoglobulin superfamily domains
antigen processing and presentation
bicellular tight junction assembly
cellular response to cadmium ion
endonucleolytic cleavage in ITS1 to separate SSU-rRNA from 5.8S-rRNA from tricistronic rRNA transcript (SSU-rRNA, 5.8S-rRNA, LSU-rRNA)
G-protein coupled receptor signaling pathway
intrinsic apoptotic signaling pathway in response to DNA damage
maturation of SSU-rRNA from tricistronic rRNA transcript (SSU-rRNA, 5.8S-rRNA, LSU-rRNA)
negative regulation of immune response
negative regulation of neuron apoptotic process
negative regulation of nucleobase-containing compound metabolic process
nucleobase-containing compound transport
positive regulation of aspartic-type endopeptidase activity involved in amyloid precursor protein catabolic process
positive regulation of cell-cell adhesion
positive regulation of cell-matrix adhesion
positive regulation of cellular component organization
positive regulation of cellular protein metabolic process
positive regulation of cellular response to growth factor stimulus
positive regulation of cellular response to hypoxia
positive regulation of cellular response to transforming growth factor beta stimulus
positive regulation of cellular response to vascular endothelial growth factor stimulus
positive regulation of cellular response to vitamin D
positive regulation of cellular response to vitamin D
positive regulation of cellular response to vitamin D