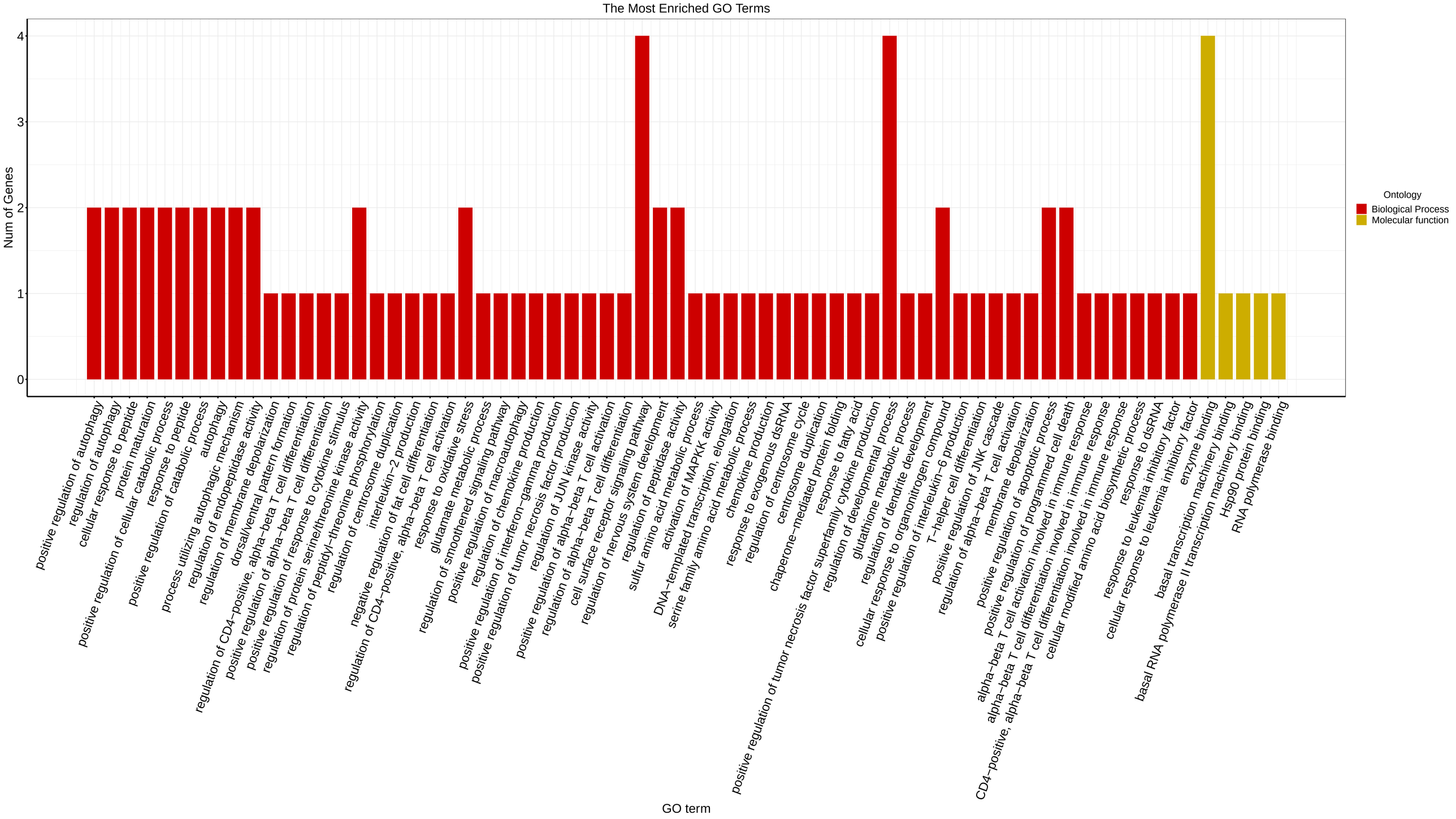
The Most Enriched GO Terms
Num of Genes
0
1
2
3
4
GO term
positive regulation of autophagy
regulation of autophagy
cellular response to peptide
protein maturation
positive regulation of cellular catabolic process
response to peptide
positive regulation of catabolic process
autophagy
process utilizing autophagic mechanism
regulation of endopeptidase activity
regulation of membrane depolarization
dorsal/ventral pattern formation
regulation of CD4−positive, alpha−beta T cell differentiation
positive regulation of alpha−beta T cell differentiation
regulation of response to cytokine stimulus
regulation of protein serine/threonine kinase activity
regulation of peptidyl−threonine phosphorylation
negative regulation of centrosome duplication
regulation of CD4−positive, alpha−beta T cell differentiation
interleukin−2 production
response to oxidative stress
glutamate metabolic process
regulation of smoothened signaling pathway
positive regulation of macroautophagy
regulation of chemokine production
positive regulation of interferon−gamma production
positive regulation of tumor necrosis factor production
regulation of JUN kinase activity
positive regulation of alpha−beta T cell activation
regulation of alpha−beta T cell differentiation
cell surface receptor signaling pathway
regulation of nervous system development
regulation of peptidase activity
sulfur amino acid metabolic process
activation of MAPKK activity
DNA−templated transcription, elongation
serine family amino acid metabolic process
chemokine production
response to exogenous dsRNA
regulation of centrosome cycle
centrosome duplication
chaperone−mediated protein folding
response to fatty acid
positive regulation of tumor necrosis factor superfamily cytokine production
regulation of developmental process
glutathione metabolic process
regulation of dendrite development
cellular response to organonitrogen compound
positive regulation of interleukin−6 production
T−helper cell differentiation
positive regulation of JNK cascade
regulation of alpha−beta T cell activation
membrane depolarization
positive regulation of apoptotic process
alpha−beta T cell activation involved in immune response
positive regulation of programmed cell death
alpha−beta T cell differentiation involved in immune response
CD4−positive, alpha−beta T cell differentiation involved in immune response
cellular modified amino acid biosynthetic process
response to leukemia inhibitory factor
response to dsRNA
cellular response to leukemia inhibitory factor
enzyme binding
basal transcription machinery binding
basal RNA polymerase II transcription machinery binding
Hsp90 protein binding
RNA polymerase binding
Ontology
Biological Process
Molecular function