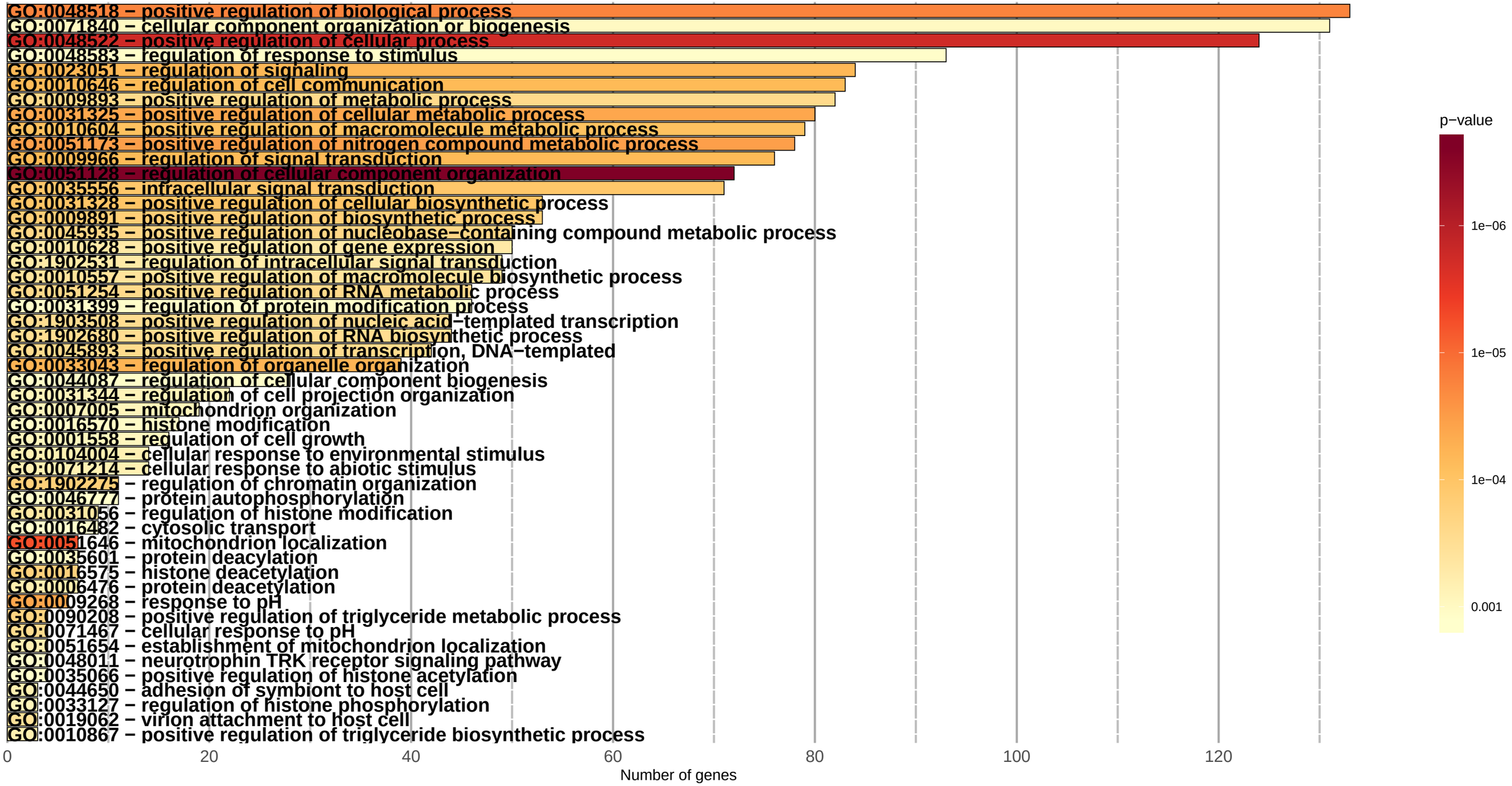
GO:0048518 − positive regulation of biological process
GO:0071840 − cellular component organization or biogenesis
GO:0048522 − positive regulation of cellular process
GO:0048583 − regulation of response to stimulus
GO:0023051 − regulation of signaling
GO:0010646 − regulation of cell communication
GO:0009893 − positive regulation of metabolic process
GO:0031325 − positive regulation of cellular metabolic process
GO:0010604 − positive regulation of macromolecule metabolic process
GO:0051173 − positive regulation of nitrogen compound metabolic process
GO:0009966 − regulation of signal transduction
GO:0051128 − regulation of cellular component organization
GO:0035556 − intracellular signal transduction
GO:0031328 − positive regulation of cellular biosynthetic process
GO:0009891 − positive regulation of biosynthetic process
GO:0045935 − positive regulation of nucleobase−containing compound metabolic process
GO:0010628 − positive regulation of gene expression
GO:1902531 − regulation of intracellular signal transduction
GO:0010557 − positive regulation of macromolecule biosynthetic process
GO:0051254 − positive regulation of RNA metabolic process
GO:0031399 − regulation of protein modification process
GO:1903508 − positive regulation of nucleic acid−templated transcription
GO:1902680 − positive regulation of RNA biosynthetic process
GO:0045893 − positive regulation of transcription, DNA−templated
GO:0033043 − regulation of organelle organization
GO:0044087 − regulation of cellular component biogenesis
GO:0031344 − regulation of cell projection organization
GO:0007005 − mitochondrion organization
GO:0016570 − histone modification
GO:0001558 − regulation of cell growth
GO:0104004 − cellular response to environmental stimulus
GO:0071214 − cellular response to abiotic stimulus
GO:1902275 − regulation of chromatin organization
GO:0046777 − protein autophosphorylation
GO:0031056 − regulation of histone modification
GO:0016482 − cytosolic transport
GO:0051646 − mitochondrion localization
GO:0035601 − protein deacylation
GO:0016575 − histone deacetylation
GO:0006476 − protein deacetylation
GO:0009268 − response to pH
GO:0090208 − positive regulation of triglyceride metabolic process
GO:0071467 − cellular response to pH
GO:0051654 − establishment of mitochondrion localization
GO:0048011 − neurotrophin TRK receptor signaling pathway
GO:0035066 − positive regulation of histone acetylation
GO:0044650 − adhesion of symbiont to host cell
GO:0033127 − regulation of histone phosphorylation
GO:0019062 − virion attachment to host cell
GO:0010867 − positive regulation of triglyceride biosynthetic process
0
20
40
60
80
100
120
Number of genes
p−value
1e−06
1e−05
1e−04
0.001