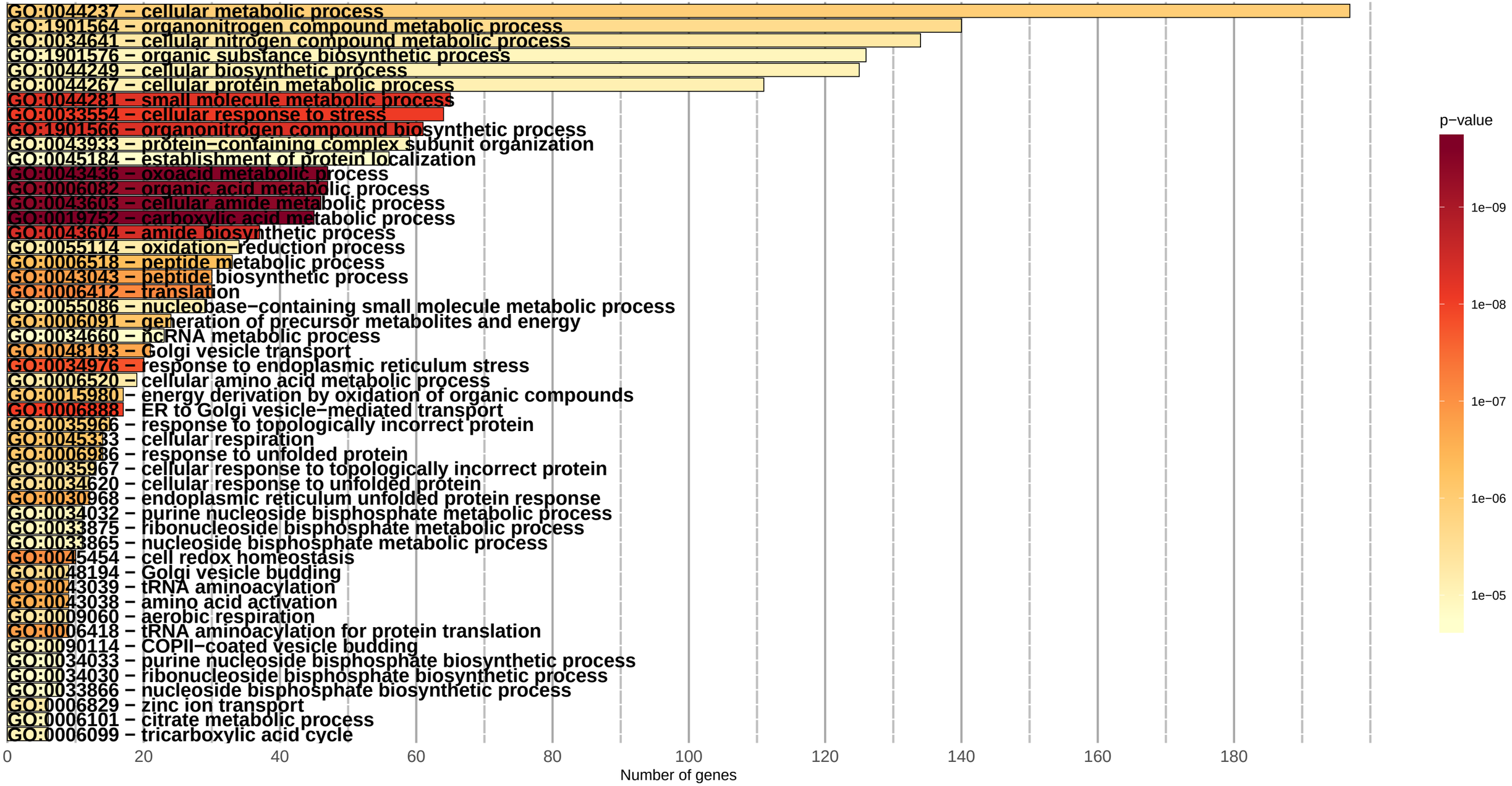
GO:0044237 − cellular metabolic process
GO:1901564 − organonitrogen compound metabolic process
GO:0034641 − cellular nitrogen compound metabolic process
GO:1901576 − organic substance biosynthetic process
GO:0044249 − cellular biosynthetic process
GO:0044267 − cellular protein metabolic process
GO:0044281 − small molecule metabolic process
GO:0033554 − cellular response to stress
GO:1901566 − organonitrogen compound biosynthetic process
GO:0043933 − protein−containing complex subunit organization
GO:0045184 − establishment of protein localization
GO:0043436 − oxoacid metabolic process
GO:0006082 − organic acid metabolic process
GO:0043603 − cellular amide metabolic process
GO:0019752 − carboxylic acid metabolic process
GO:0043604 − amide biosynthetic process
GO:0055114 − oxidation−reduction process
GO:0006518 − peptide metabolic process
GO:0043043 − peptide biosynthetic process
GO:0006412 − translation
GO:0055086 − nucleobase−containing small molecule metabolic process
GO:0006091 − generation of precursor metabolites and energy
GO:0034660 − ncRNA metabolic process
GO:0048193 − Golgi vesicle transport
GO:0034976 − response to endoplasmic reticulum stress
GO:0006520 − cellular amino acid metabolic process
GO:0015980 − energy derivation by oxidation of organic compounds
GO:0006888 − ER to Golgi vesicle−mediated transport
GO:0035966 − response to topologically incorrect protein
GO:0045333 − cellular respiration
GO:0006986 − response to unfolded protein
GO:0035967 − cellular response to topologically incorrect protein
GO:0034620 − cellular response to unfolded protein
GO:0030968 − endoplasmic reticulum unfolded protein response
GO:0034032 − purine nucleoside bisphosphate metabolic process
GO:0033875 − ribonucleoside bisphosphate metabolic process
GO:0033865 − nucleoside bisphosphate metabolic process
GO:0045454 − cell redox homeostasis
GO:0048194 − Golgi vesicle budding
GO:0043039 − tRNA aminoacylation
GO:0043038 − amino acid activation
GO:0009060 − aerobic respiration
GO:0006418 − tRNA aminoacylation for protein translation
GO:0090114 − COPII−coated vesicle budding
GO:0034033 − purine nucleoside bisphosphate biosynthetic process
GO:0034030 − ribonucleoside bisphosphate biosynthetic process
GO:0033866 − nucleoside bisphosphate biosynthetic process
GO:0006829 − zinc ion transport
GO:0006101 − citrate metabolic process
GO:0006099 − tricarboxylic acid cycle
0
20
40
60
80
100
120
140
160
180
Number of genes
p−value
1e−09
1e−08
1e−07
1e−06
1e−05