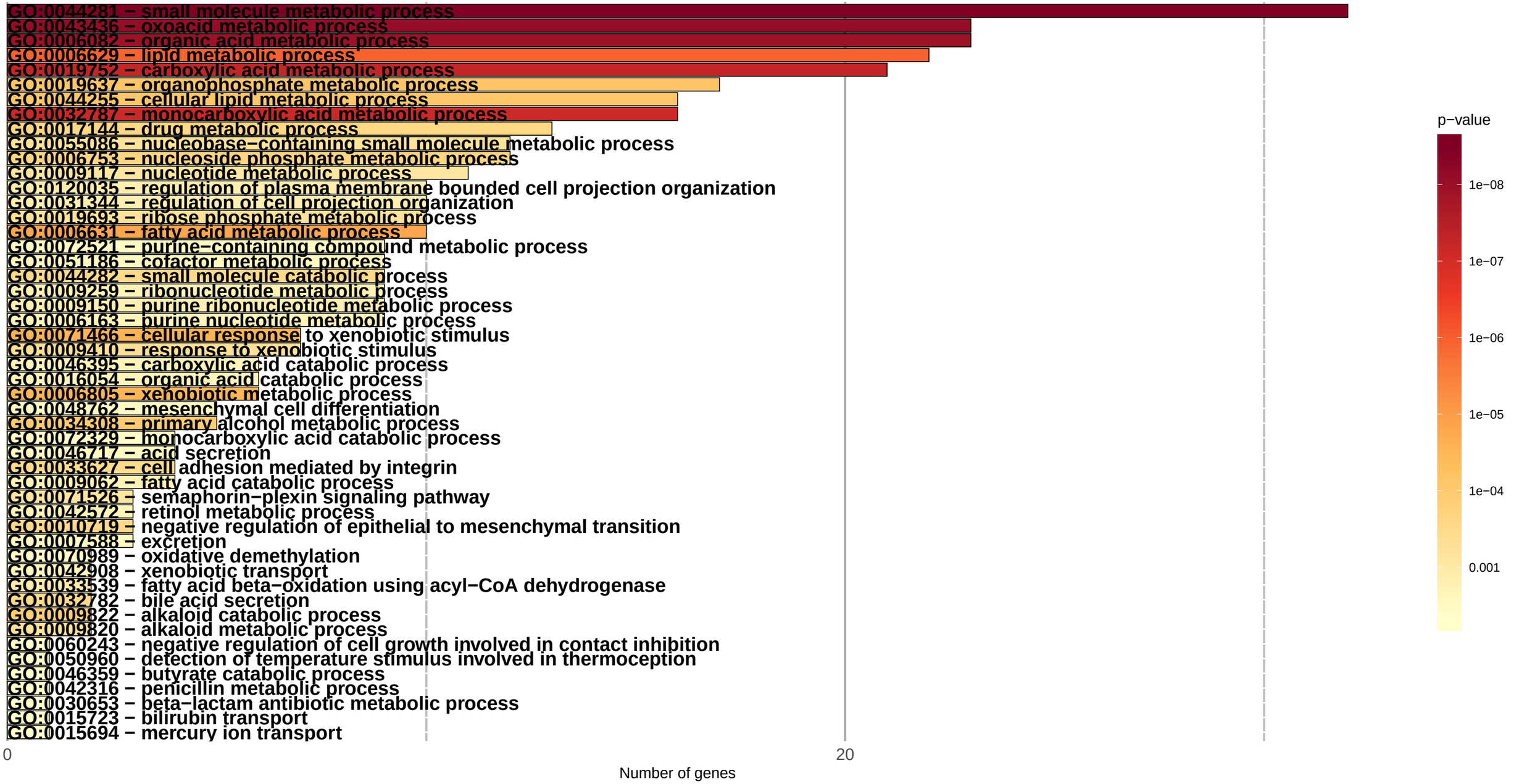
GO:0044281 − small molecule metabolic process
GO:0043436 − oxoacid metabolic process
GO:0006082 − organic acid metabolic process
GO:0006629 − lipid metabolic process
GO:0019752 − carboxylic acid metabolic process
GO:0019637 − organophosphate metabolic process
GO:0044255 − cellular lipid metabolic process
GO:0032787 − monocarboxylic acid metabolic process
GO:0017144 − drug metabolic process
GO:0055086 − nucleobase−containing small molecule metabolic process
GO:0006753 − nucleoside phosphate metabolic process
GO:0009117 − nucleotide metabolic process
GO:0120035 − regulation of plasma membrane bounded cell projection organization
GO:0031344 − regulation of cell projection organization
GO:0019693 − ribose phosphate metabolic process
GO:0006631 − fatty acid metabolic process
GO:0072521 − purine−containing compound metabolic process
GO:0051186 − cofactor metabolic process
GO:0044282 − small molecule catabolic process
GO:0009259 − ribonucleotide metabolic process
GO:0009150 − purine ribonucleotide metabolic process
GO:0006163 − purine nucleotide metabolic process
GO:0071466 − cellular response to xenobiotic stimulus
GO:0009410 − response to xenobiotic stimulus
GO:0046395 − carboxylic acid catabolic process
GO:0016054 − organic acid catabolic process
GO:0006805 − xenobiotic metabolic process
GO:0048762 − mesenchymal cell differentiation
GO:0034308 − primary alcohol metabolic process
GO:0072329 − monocarboxylic acid catabolic process
GO:0046717 − acid secretion
GO:0033627 − cell adhesion mediated by integrin
GO:0009062 − fatty acid catabolic process
GO:0071526 − semaphorin−plexin signaling pathway
GO:0042572 − retinol metabolic process
GO:0010719 − negative regulation of epithelial to mesenchymal transition
GO:0007588 − excretion
GO:0070989 − oxidative demethylation
GO:0042908 − xenobiotic transport
GO:0033539 − fatty acid beta−oxidation using acyl−CoA dehydrogenase
GO:0032782 − bile acid secretion
GO:0009822 − alkaloid catabolic process
GO:0009820 − alkaloid metabolic process
GO:0060243 − negative regulation of cell growth involved in contact inhibition
GO:0050960 − detection of temperature stimulus involved in thermoception
GO:0046359 − butyrate catabolic process
GO:0042316 − penicillin metabolic process
GO:0030653 − beta−lactam antibiotic metabolic process
GO:0015723 − bilirubin transport
GO:0015694 − mercury ion transport
0
20
Number of genes
p−value
1e−08
1e−07
1e−06
1e−05
1e−04
0.001