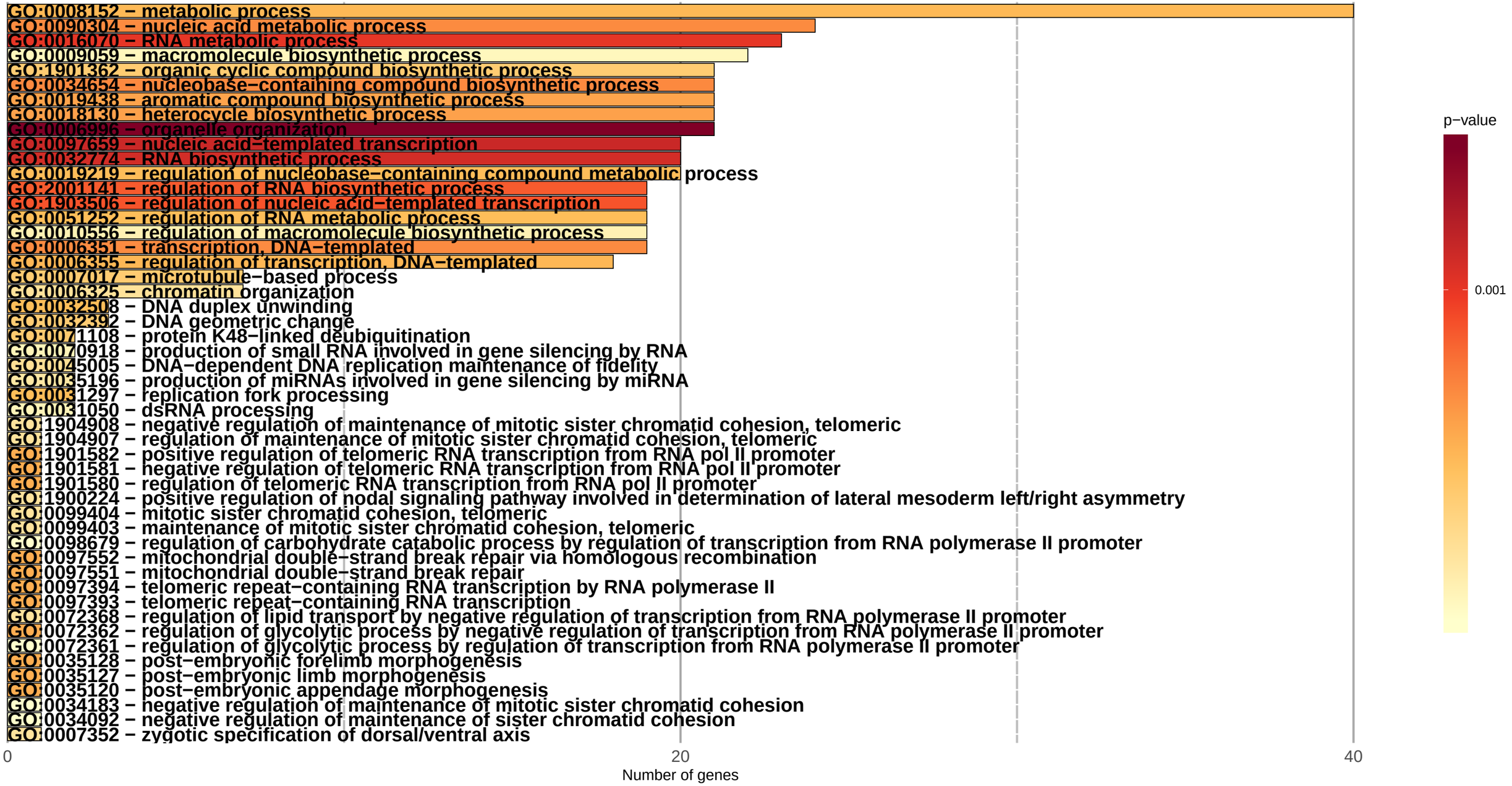
GO:0008152 − metabolic process
GO:0090304 − nucleic acid metabolic process
GO:0016070 − RNA metabolic process
GO:0009059 − macromolecule biosynthetic process
GO:1901362 − organic cyclic compound biosynthetic process
GO:0034654 − nucleobase−containing compound biosynthetic process
GO:0019438 − aromatic compound biosynthetic process
GO:0018130 − heterocycle biosynthetic process
GO:0006996 − organelle organization
GO:0097659 − nucleic acid−templated transcription
GO:0032774 − RNA biosynthetic process
GO:0019219 − regulation of nucleobase−containing compound metabolic process
GO:2001141 − regulation of RNA biosynthetic process
GO:1903506 − regulation of nucleic acid−templated transcription
GO:0051252 − regulation of RNA metabolic process
GO:0010556 − regulation of macromolecule biosynthetic process
GO:0006351 − transcription, DNA−templated
GO:0006355 − regulation of transcription, DNA−templated
GO:0007017 − microtubule−based process
GO:0006325 − chromatin organization
GO:0032508 − DNA duplex unwinding
GO:0032392 − DNA geometric change
GO:0071108 − protein K48−linked deubiquitination
GO:0070918 − production of small RNA involved in gene silencing by RNA
GO:0045005 − DNA−dependent DNA replication maintenance of fidelity
GO:0035196 − production of miRNAs involved in gene silencing by miRNA
GO:0031297 − replication fork processing
GO:0031050 − dsRNA processing
GO:1904908 − negative regulation of maintenance of mitotic sister chromatid cohesion, telomeric
GO:1904907 − regulation of maintenance of mitotic sister chromatid cohesion, telomeric
GO:1901582 − positive regulation of telomeric RNA transcription from RNA pol II promoter
GO:1901581 − negative regulation of telomeric RNA transcription from RNA pol II promoter
GO:1901580 − regulation of telomeric RNA transcription from RNA pol II promoter
GO:1900224 − positive regulation of nodal signaling pathway involved in determination of lateral mesoderm left/right asymmetry
GO:0099404 − mitotic sister chromatid cohesion, telomeric
GO:0099403 − maintenance of mitotic sister chromatid cohesion, telomeric
GO:0098679 − regulation of carbohydrate catabolic process by regulation of transcription from RNA polymerase II promoter
GO:0097552 − mitochondrial double−strand break repair via homologous recombination
GO:0097551 − mitochondrial double−strand break repair
GO:0097394 − telomeric repeat−containing RNA transcription by RNA polymerase II
GO:0097393 − telomeric repeat−containing RNA transcription
GO:0072368 − regulation of lipid transport by negative regulation of transcription from RNA polymerase II promoter
GO:0072362 − regulation of glycolytic process by negative regulation of transcription from RNA polymerase II promoter
GO:0072361 − regulation of glycolytic process by regulation of transcription from RNA polymerase II promoter
GO:0035128 − post−embryonic forelimb morphogenesis
GO:0035127 − post−embryonic limb morphogenesis
GO:0035120 − post−embryonic appendage morphogenesis
GO:0034183 − negative regulation of maintenance of mitotic sister chromatid cohesion
GO:0034092 − negative regulation of maintenance of sister chromatid cohesion
GO:0007352 − zygotic specification of dorsal/ventral axis
0
20
40
Number of genes
p−value
0.001