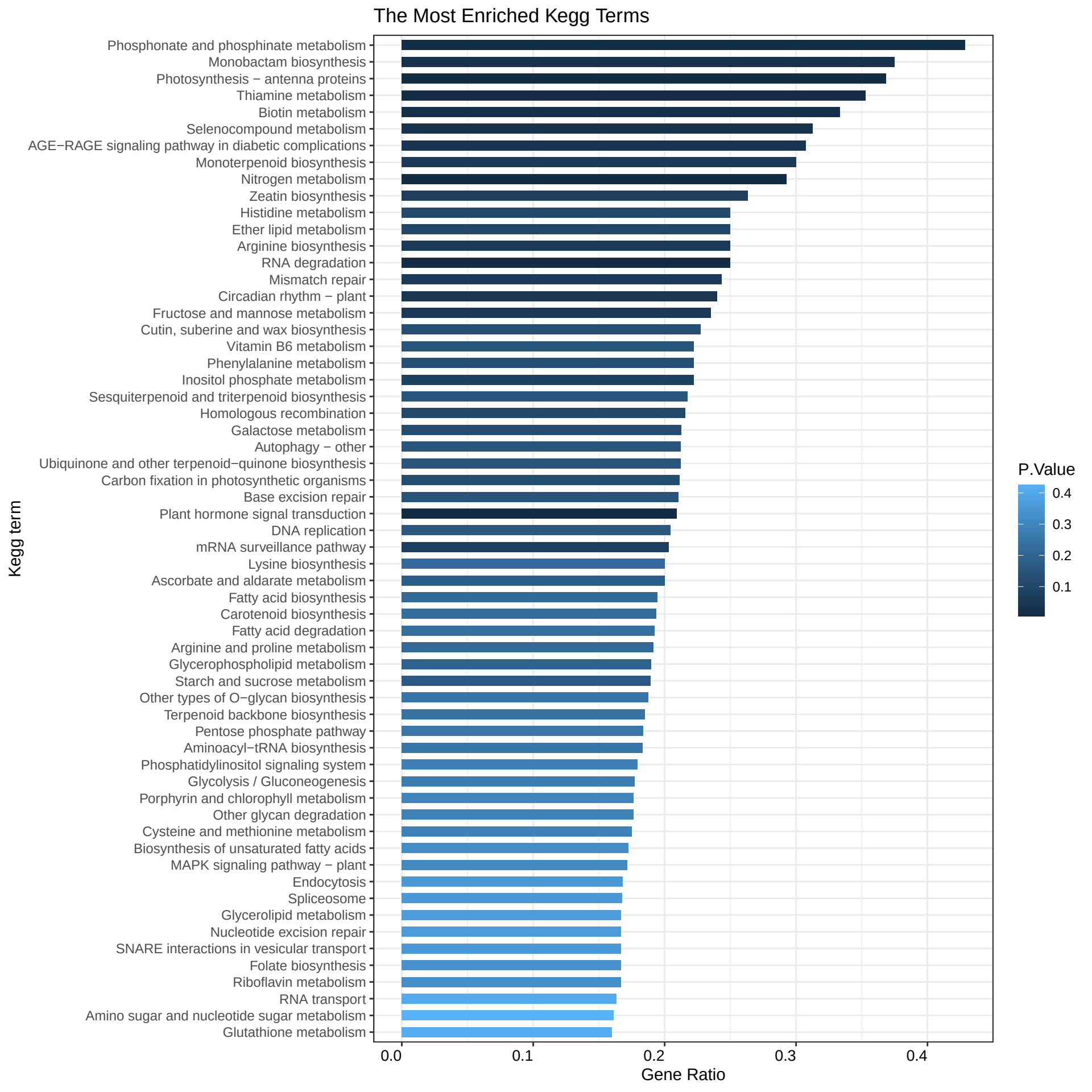
The Most Enriched Kegg Terms
Phosphonate and phosphinate metabolism
Monobactam biosynthesis
Photosynthesis − antenna proteins
Thiamine metabolism
Biotin metabolism
Selenocompound metabolism
AGE−RAGE signaling pathway in diabetic complications
Monoterpenoid biosynthesis
Nitrogen metabolism
Zeatin biosynthesis
Histidine metabolism
Ether lipid metabolism
Arginine biosynthesis
RNA degradation
Mismatch repair
Circadian rhythm − plant
Fructose and mannose metabolism
Cutin, suberine and wax biosynthesis
Vitamin B6 metabolism
Phenylalanine metabolism
Inositol phosphate metabolism
Sesquiterpenoid and triterpenoid biosynthesis
Homologous recombination
Galactose metabolism
Autophagy − other
Ubiquinone and other terpenoid−quinone biosynthesis
Carbon fixation in photosynthetic organisms
Base excision repair
Plant hormone signal transduction
DNA replication
mRNA surveillance pathway
Lysine biosynthesis
Ascorbate and aldarate metabolism
Fatty acid biosynthesis
Carotenoid biosynthesis
Fatty acid degradation
Arginine and proline metabolism
Glycerophospholipid metabolism
Starch and sucrose metabolism
Other types of O−glycan biosynthesis
Terpenoid backbone biosynthesis
Pentose phosphate pathway
Aminoacyl−tRNA biosynthesis
Phosphatidylinositol signaling system
Glycolysis / Gluconeogenesis
Porphyrin and chlorophyll metabolism
Other glycan degradation
Cysteine and methionine metabolism
Biosynthesis of unsaturated fatty acids
MAPK signaling pathway − plant
Endocytosis
Spliceosome
Glycerolipid metabolism
Nucleotide excision repair
SNARE interactions in vesicular transport
Folate biosynthesis
Riboflavin metabolism
RNA transport
Amino sugar and nucleotide sugar metabolism
Glutathione metabolism
Kegg term
0.0
0.1
0.2
0.3
0.4
Gene Ratio
P.Value
0.4
0.3
0.2
0.1