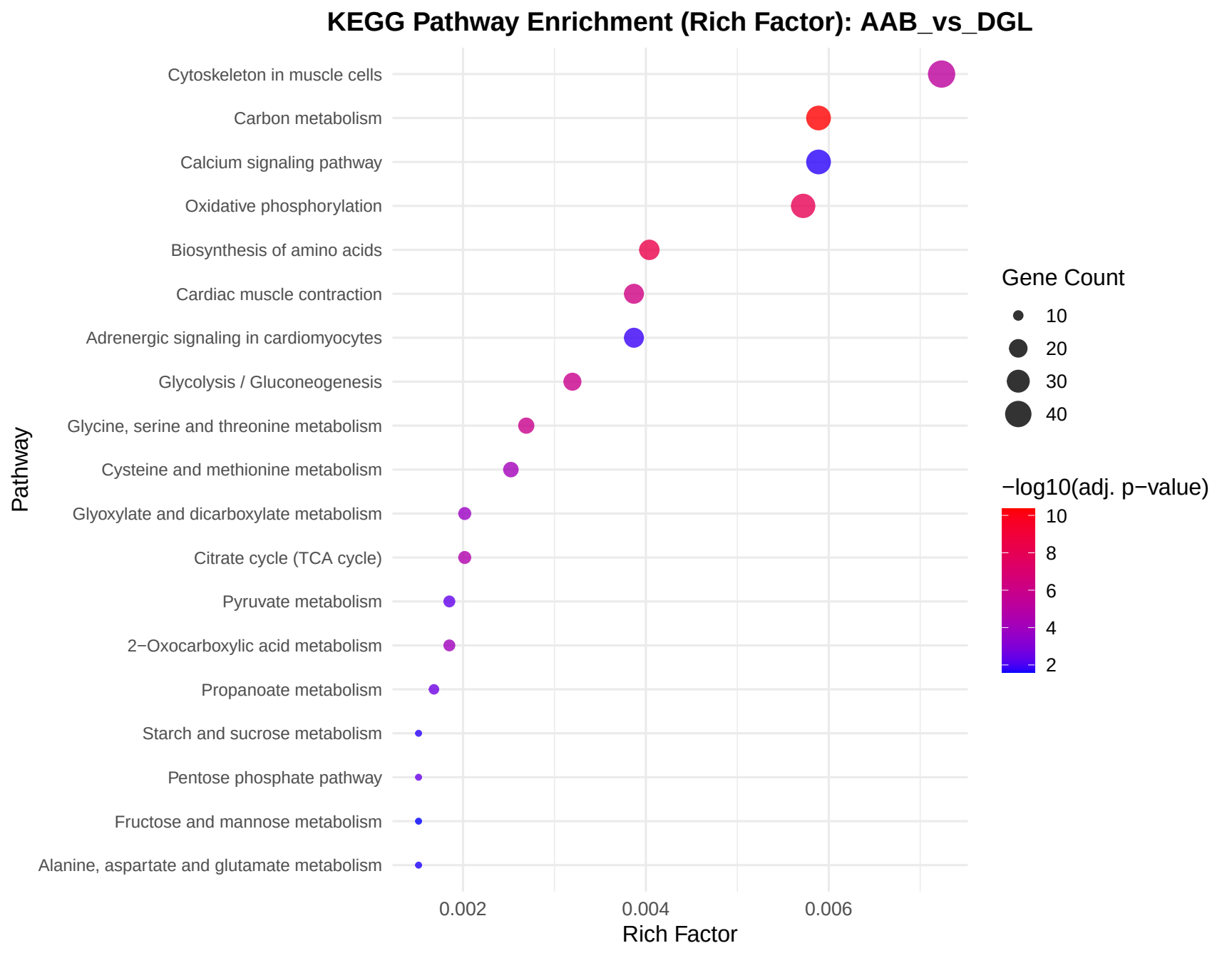
KEGG Pathway Enrichment (Rich Factor): AAB_vs_DGL
Cytoskeleton in muscle cells
Carbon metabolism
Calcium signaling pathway
Oxidative phosphorylation
Biosynthesis of amino acids
Cardiac muscle contraction
Adrenergic signaling in cardiomyocytes
Glycolysis / Gluconeogenesis
Glycine, serine and threonine metabolism
Cysteine and methionine metabolism
Glyoxylate and dicarboxylate metabolism
Citrate cycle (TCA cycle)
Pyruvate metabolism
2−Oxocarboxylic acid metabolism
Propanoate metabolism
Starch and sucrose metabolism
Pentose phosphate pathway
Fructose and mannose metabolism
Alanine, aspartate and glutamate metabolism
Pathway
0.002
0.004
0.006
Rich Factor
Gene Count
10
20
30
40
−log10(adj. p−value)
10
8
6
4
2