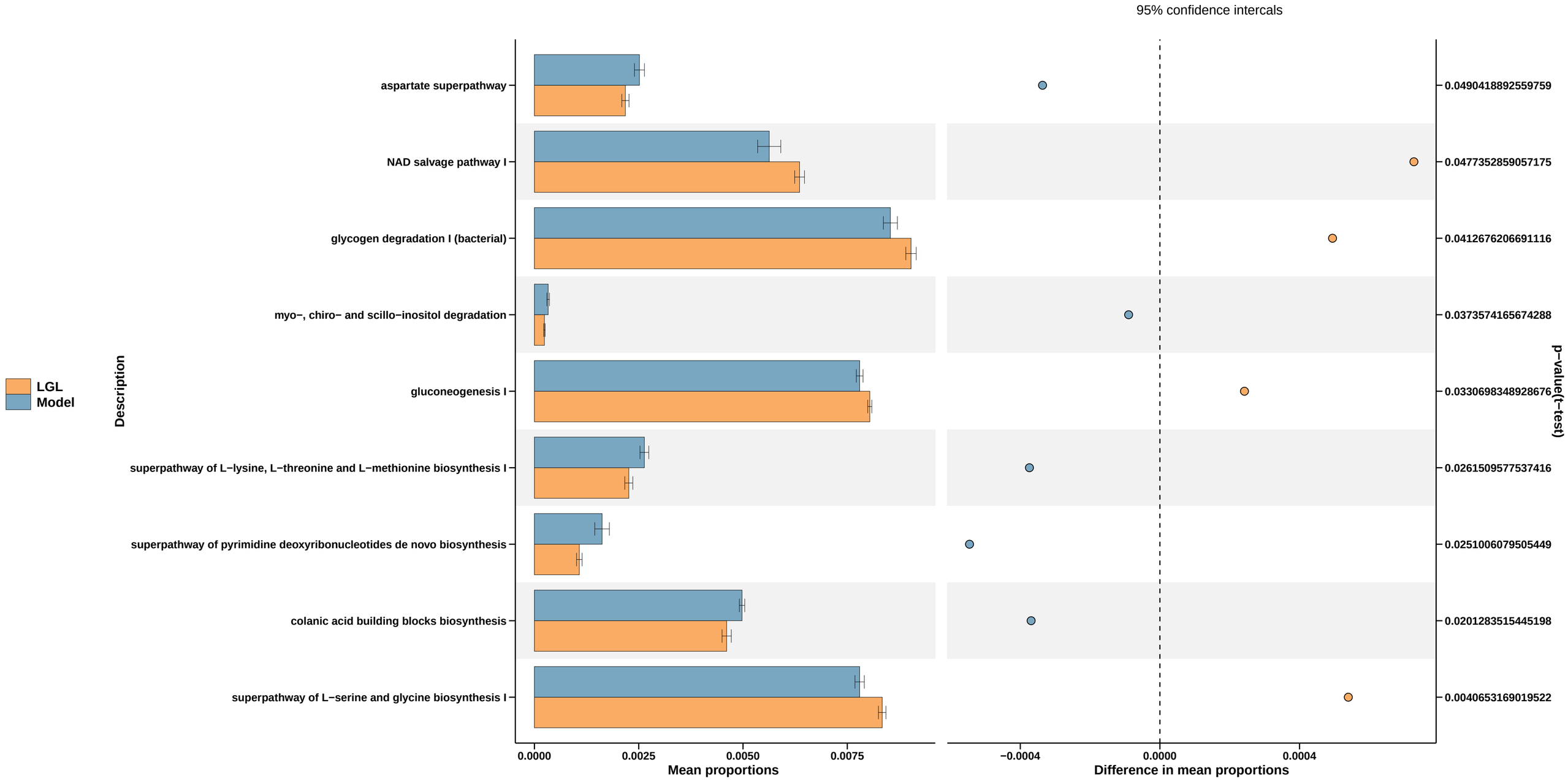
95% confidence intercals
LGL
Model
Description
aspartate superpathway
NAD salvage pathway I
glycogen degradation I (bacterial)
myo−, chiro− and scillo−inositol degradation
gluconeogenesis I
superpathway of L−lysine, L−threonine and L−methionine biosynthesis I
superpathway of pyrimidine deoxyribonucleotides de novo biosynthesis
colanic acid building blocks biosynthesis
superpathway of L−serine and glycine biosynthesis I
0.0000
0.0025
0.0050
0.0075
Mean proportions
−0.0004
0.0000
0.0004
Difference in mean proportions
p−value(t−test)
0.0490418892559759
0.0477352859057175
0.0412676206691116
0.0373574165674288
0.0330698348928676
0.0261509577537416
0.0251006079505449
0.0201283515445198
0.0040653169019522