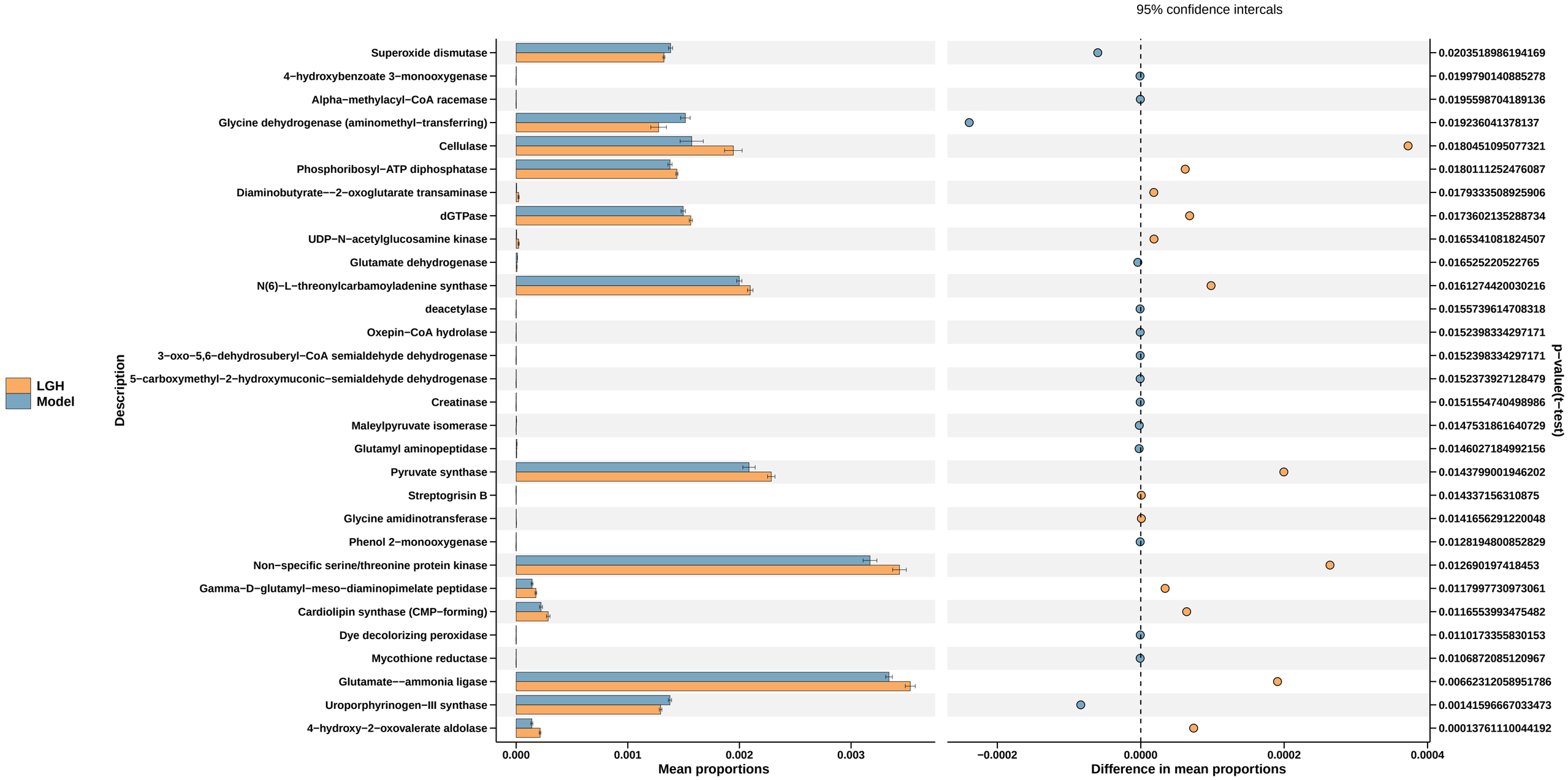
95% confidence intercals
LGH
Model
Description
Superoxide dismutase
4−hydroxybenzoate 3−monooxygenase
Alpha−methylacyl−CoA racemase
Glycine dehydrogenase (aminomethyl−transferring)
Cellulase
Phosphoribosyl−ATP diphosphatase
Diaminobutyrate−−2−oxoglutarate transaminase
dGTPase
UDP−N−acetylglucosamine kinase
Glutamate dehydrogenase
N(6)−L−threonylcarbamoyladenine synthase
deacetylase
Oxepin−CoA hydrolase
3−oxo−5,6−dehydrosuberyl−CoA semialdehyde dehydrogenase
5−carboxymethyl−2−hydroxymuconic−semialdehyde dehydrogenase
Creatinase
Maleylpyruvate isomerase
Glutamyl aminopeptidase
Pyruvate synthase
Streptogrisin B
Glycine amidinotransferase
Phenol 2−monooxygenase
Non−specific serine/threonine protein kinase
Gamma−D−glutamyl−meso−diaminopimelate peptidase
Cardiolipin synthase (CMP−forming)
Dye decolorizing peroxidase
Mycothione reductase
Glutamate−−ammonia ligase
Uroporphyrinogen−III synthase
4−hydroxy−2−oxovalerate aldolase
0.000
0.001
0.002
0.003
Mean proportions
−0.0002
0.0000
0.0002
0.0004
Difference in mean proportions
p−value(t−test)
0.0203518986194169
0.0199790140885278
0.0195598704189136
0.019236041378137
0.0180451095077321
0.0180111252476087
0.0179333508925906
0.0173602135288734
0.0165341081824507
0.016525220522765
0.0161274420030216
0.0155739614708318
0.0152398334297171
0.0152398334297171
0.0152373927128479
0.0151554740498986
0.0147531861640729
0.0146027184992156
0.0143799001946202
0.014337156310875
0.0141656291220048
0.0128194800852829
0.012690197418453
0.0117997730973061
0.0116553993475482
0.0110173355830153
0.0106872085120967
0.00662312058951786
0.00141596667033473
0.00013761110044192