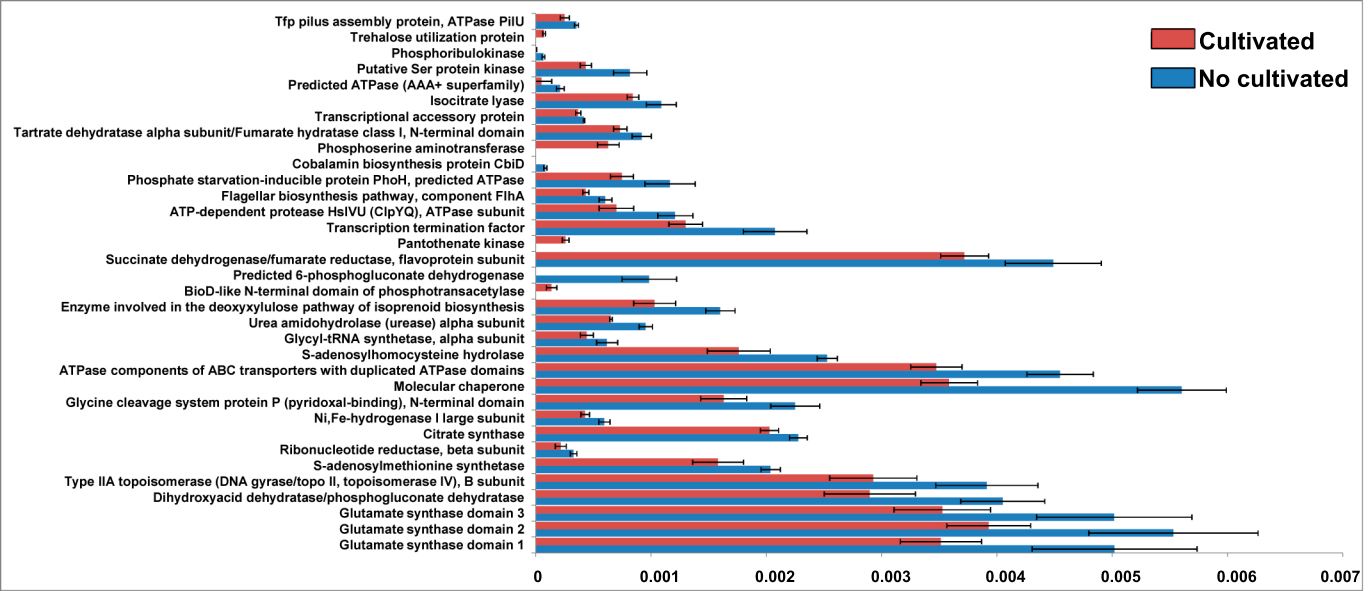
Cultivated
No cultivated
Tfp pilus assembly protein, ATPase PilU
Trehalose utilization protein
Phosphoribulokinase
Putative Ser protein kinase
Predicted ATPase (AAA+ superfamily)
Isocitrate lyase
Transcriptional accessory protein
Tartrate dehydratase alpha subunit/Fumarate hydratase class I, N-terminal domain
Phosphoserine aminotransferase
Cobalamin biosynthesis protein CbiD
Phosphate starvation-inducible protein PhoH, predicted ATPase
Flagellar biosynthesis pathway, component FlhA
ATP-dependent protease HslVU (ClpYQ), ATPase subunit
Transcription termination factor
Pantothenate kinase
Succinate dehydrogenase/fumarate reductase, flavoprotein subunit
Predicted 6-phosphogluconate dehydrogenase
BioD-like N-terminal domain of phosphotransacetylase
Enzyme involved in the deoxyxylulose pathway of isoprenoid biosynthesis
Urea amidohydrolase (urease) alpha subunit
Glycyl-tRNA synthetase, alpha subunit
S-adenosylhomocysteine hydrolase
ATPase components of ABC transporters with duplicated ATPase domains
Molecular chaperone
Glycine cleavage system protein P (pyridoxal-binding), N-terminal domain
Ni,Fe-hydrogenase I large subunit
Citrate synthase
Ribonucleotide reductase, beta subunit
S-adenosylmethionine synthetase
Type IIA topoisomerase (DNA gyrase/topo II, topoisomerase IV), B subunit
Dihydroxyacid dehydratase/phosphogluconate dehydratase
Glutamate synthase domain 3
Glutamate synthase domain 2
Glutamate synthase domain 1
0
0.001
0.002
0.003
0.004
0.005
0.006
0.007