Blue_vs_White

Expression
- 100
- 200
- 300
- 400
- 500
- 600

Cluster
- Cluster 1
- Cluster 2
- Cluster 3
- Cluster 4
- Cluster 5
- Cluster 6
- Cluster 7

single−strand DNA−binding protein

preprotein translocase subunit YajC

ribulose−bisphosphate carboxylase small chain [EC:4.1.1.39]

DNA−binding protein HU−beta

histidyl−tRNA synthetase [EC:6.1.1.21]

aldehyde dehydrogenase [EC:1.2.1.−]

ribose transport system ATP−binding protein [EC:7.5.2.7]

3−hydroxybutyrate dehydrogenase [EC:1.1.1.30]

reactive chlorine resistance protein C

conjugal transfer pilin signal peptidase Trbl hydroxymethylglutaryl−CoA reductase [EC:1.1.1.88]

uncharacterized protein involved in response to NO

large subunit ribosomal protein L1 ribosomal protein S20

aspartyl−tRNA(Asn)/glutamyl−tRNA(Gln) amidotransferase subunit C [EC:6.3.5.6 6.3.5.7]

biotin transport system ATP−binding protein [EC:7.6.2.−]

small subunit ribosomal protein S16

putative acetyl−hydrolase [EC:2.6.1.2] ammonium transporter

2−hydrolase [EC:2.6.1.2] 4−oxalocrotonate tautomerase [EC:5.3.2.6]

cell division protein ZapE

glutathione S−transferase [EC:2.5.1.18]

2−oxoglutarate dehydrogenase E1 component [EC:1.2.4.2]

purine−nucleoside phosphorylase [EC:2.4.2.1]

NFU1 iron−sulfur cluster scaffold homolog, mitochondrial

acetoacetyl−CoA reductase [EC:1.1.1.36] GTPase

P−value

Fold change