GO_Biological_Process_2021

Horizontal bar chart titled "GO_Biological_Process_2021" with y-axis labeled "Term" and x-axis labeled "Enrichment log(combined score)". Bars from top to bottom:

- L−serine biosynthetic process (GO:0006564)
- aerobic electron transport chain (GO:0019646)
- mitochondrial ATP synthesis coupled electron transport (GO:0042775)
- L−serine metabolic process (GO:0006563)
- cellular biogenic amine catabolic process (GO:0042402)
- serine family amino acid biosynthetic process (GO:0009070)
- mitochondrial electron transport, ubiquinol to cytochrome c (GO:0006122)
- mitochondrial electron transport, NADH to ubiquinone (GO:0006120)
- substantia nigra development (GO:0021762)
- regulation of adherens junction organization (GO:1903391)
- protein lipoylation (GO:0009249)
- negative regulation of lipid transport (GO:0032369)
- glutamate catabolic process (GO:0006538)
- endothelin receptor signaling pathway (GO:0086100)
- cellular response to brain−derived neurotrophic factor stimulus (GO:1990416)

GO_Cellular_Component_2021

respiratory chain complex I (GO:0045271)

mitochondrial respiratory chain complex I (GO:0005747)

smooth endoplasmic reticulum membrane (GO:0030868)

U4 snRNP (GO:0005687)

mitochondrial respiratory chain complex III (GO:0005750)

mitochondrial inner membrane (GO:0005743)

organelle inner membrane (GO:0019866)

Swr1 complex (GO:0000812)

integral component of peroxisomal membrane (GO:0005779)

mitochondrial membrane (GO:0031966)

Term

Enrichment log(combined score)

0   2   4

GO_Molecular_Function_2021

Bar chart titled "GO_Molecular_Function_2021" with x-axis "Enrichment log(combined score)" and y-axis "Term".

- ubiquinol−cytochrome−c reductase activity (GO:0008121)
- oxidoreduction−driven active transmembrane transporter activity (GO:0015453)
- thiolester hydrolase activity (GO:0016790)
- pyruvate transmembrane transporter activity (GO:0050833)
- phosphotransferase activity, nitrogenous group as acceptor (GO:0016775)
- ganglioside binding (GO:0035594)
- creatine kinase activity (GO:0004111)
- fructose binding (GO:0070061)
- NADH dehydrogenase (ubiquinone) activity (GO:0008137)
- NADH dehydrogenase (quinone) activity (GO:0050136)