GO_Biological_Process_2017

endosomal transport (GO:0016197)

RNA splicing (GO:0008380)

establishment of Golgi localization (GO:0051683)

multivesicular body assembly (GO:0036258)

gluconeogenesis (GO:0006094)

translation (GO:0006412)

establishment of epithelial cell apical/basal polarity (GO:0045198)

mitochondrial ATP synthesis coupled proton transport (GO:0042776)

regulation of NIK/NF-kappaB signaling (GO:1901222)

positive regulation of ATP biosynthetic process (GO:2001171)

$-\log_{10}$ P-value

GO_Cellular_Component_2017

mitochondrion (GO:0005739)

nucleolus (GO:0005730)

ficolin-1-rich granule lumen (GO:1904813)

secretory granule lumen (GO:0034774)

endosome (GO:0005768)

small nucleolar ribonucleoprotein complex (GO:0005732)

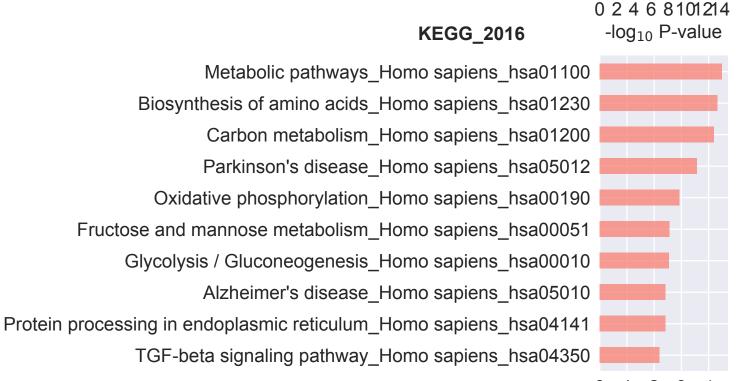catalytic step 2 spliceosome (GO:0071013)

mitochondrial inner membrane presequence translocase complex (GO:0005744)

centrosome (GO:0005813)

lysosomal membrane (GO:0005765)

$-\log_{10}$ P-value

GO_Molecular_Function_2017

RNA binding (GO:0003723)

GDP binding (GO:0019003)

myosin V binding (GO:0031489)

cadherin binding (GO:0045296)

NADH dehydrogenase activity (GO:0003954)

protein phosphatase regulator activity (GO:0019888)

protein homodimerization activity (GO:0042803)

GTP binding (GO:0005525)

proton-transporting ATP synthase activity, rotational mechanism (GO:0046933)

androgen receptor binding (GO:0050681)

$-\log_{10}$ P-value

KEGG_2016

Metabolic pathways_Homo sapiens_hsa01100

Biosynthesis of amino acids_Homo sapiens_hsa01230

Carbon metabolism_Homo sapiens_hsa01200

Parkinson's disease_Homo sapiens_hsa05012

Oxidative phosphorylation_Homo sapiens_hsa00190

Fructose and mannose metabolism_Homo sapiens_hsa00051

Glycolysis / Gluconeogenesis_Homo sapiens_hsa00010

Alzheimer's disease_Homo sapiens_hsa05010

Protein processing in endoplasmic reticulum_Homo sapiens_hsa04141

TGF-beta signaling pathway_Homo sapiens_hsa04350

$-\log_{10}$ P-value