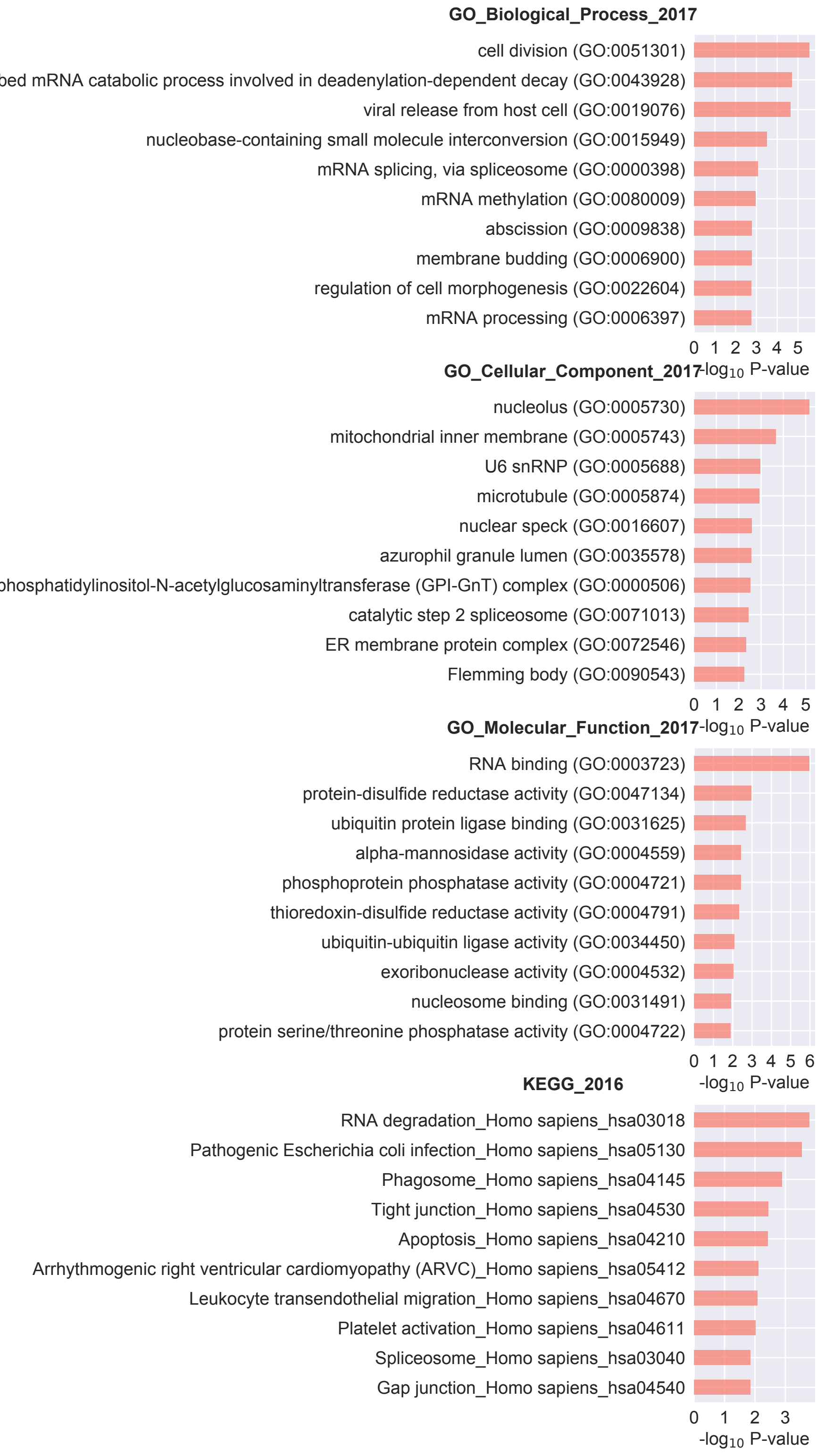
GO_Biological_Process_2017
cell division (GO:0051301)
bed mRNA catabolic process involved in deadenylation-dependent decay (GO:0043928)
viral release from host cell (GO:0019076)
nucleobase-containing small molecule interconversion (GO:0015949)
mRNA splicing, via spliceosome (GO:0000398)
mRNA methylation (GO:0080009)
abscission (GO:0009838)
membrane budding (GO:0006900)
regulation of cell morphogenesis (GO:0022604)
mRNA processing (GO:0006397)
0 1 2 3 4 5
-log10 P-value
GO_Cellular_Component_2017
nucleolus (GO:0005730)
mitochondrial inner membrane (GO:0005743)
U6 snRNP (GO:0005688)
microtubule (GO:0005874)
nuclear speck (GO:0016607)
azurophil granule lumen (GO:0035578)
phosphatidylinositol-N-acetylglucosaminyltransferase (GPI-GnT) complex (GO:0000506)
catalytic step 2 spliceosome (GO:0071013)
ER membrane protein complex (GO:0072546)
Flemming body (GO:0090543)
0 1 2 3 4 5
-log10 P-value
GO_Molecular_Function_2017
RNA binding (GO:0003723)
protein-disulfide reductase activity (GO:0047134)
ubiquitin protein ligase binding (GO:0031625)
alpha-mannosidase activity (GO:0004559)
phosphoprotein phosphatase activity (GO:0004721)
thioredoxin-disulfide reductase activity (GO:0004791)
ubiquitin-ubiquitin ligase activity (GO:0034450)
exoribonuclease activity (GO:0004532)
nucleosome binding (GO:0031491)
protein serine/threonine phosphatase activity (GO:0004722)
0 1 2 3 4 5 6
-log10 P-value
KEGG_2016
RNA degradation_Homo sapiens_hsa03018
Pathogenic Escherichia coli infection_Homo sapiens_hsa05130
Phagosome_Homo sapiens_hsa04145
Tight junction_Homo sapiens_hsa04530
Apoptosis_Homo sapiens_hsa04210
Arrhythmogenic right ventricular cardiomyopathy (ARVC)_Homo sapiens_hsa05412
Leukocyte transendothelial migration_Homo sapiens_hsa04670
Platelet activation_Homo sapiens_hsa04611
Spliceosome_Homo sapiens_hsa03040
Gap junction_Homo sapiens_hsa04540
0 1 2 3
-log10 P-value