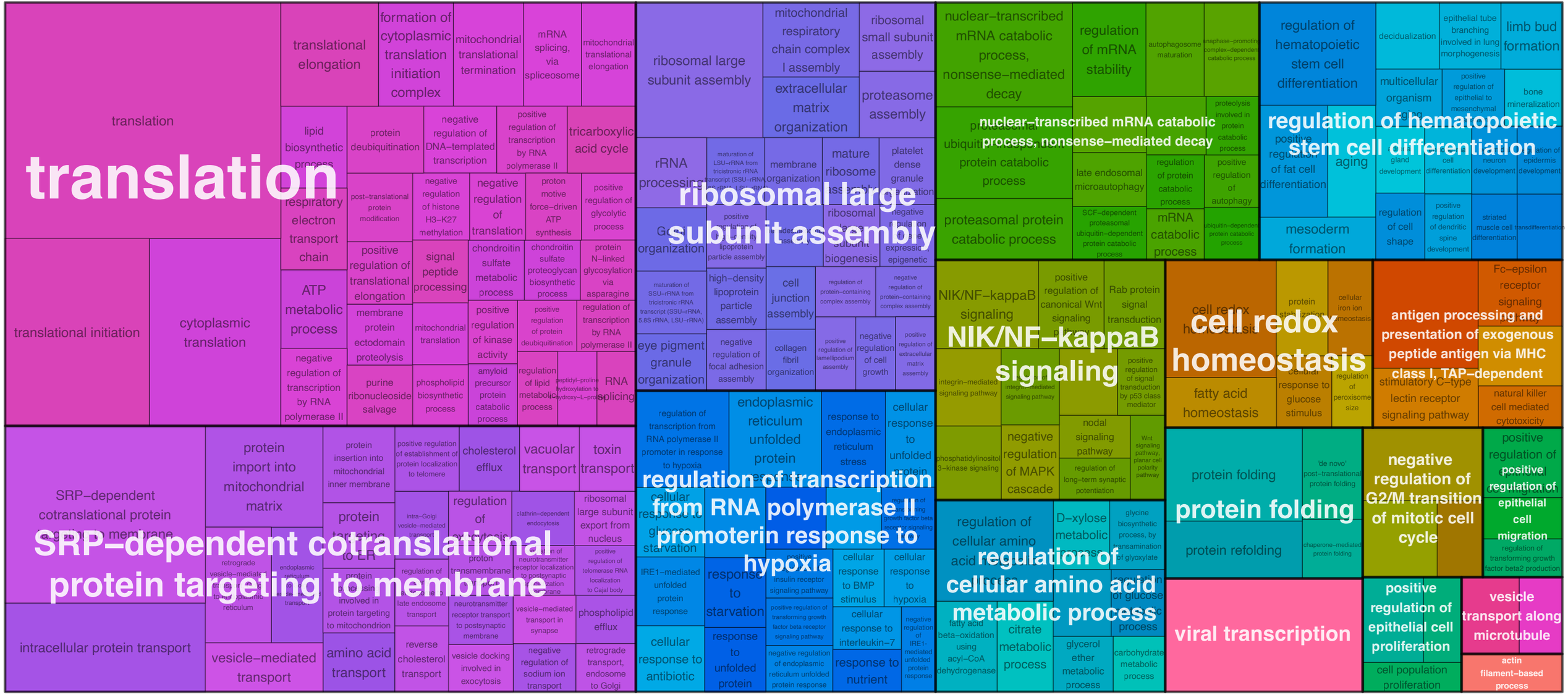
translation
translational elongation
formation of cytoplasmic translation initiation complex
mitochondrial translational termination
mRNA splicing, via spliceosome
mitochondrial translational elongation
translational initiation
cytoplasmic translation
lipid biosynthetic process
protein deubiquitination
negative regulation of DNA−templated transcription
positive regulation of transcription by RNA polymerase II
tricarboxylic acid cycle
respiratory electron transport chain
post−translational protein modification
negative regulation of histone H3−K27 methylation
negative regulation of translation
proton motive force−driven ATP synthesis
positive regulation of glycolytic process
ATP metabolic process
positive regulation of translational elongation
signal peptide processing
chondroitin sulfate metabolic process
chondroitin sulfate proteoglycan biosynthetic process
protein N−linked glycosylation via asparagine
negative regulation of transcription by RNA polymerase II
membrane protein ectodomain proteolysis
mitochondrial translation
positive regulation of kinase activity
positive regulation of protein deubiquitination
regulation of transcription by RNA polymerase II
purine ribonucleoside salvage
phospholipid biosynthetic process
amyloid precursor protein catabolic process
regulation of lipid metabolic process
peptidyl−proline hydroxylation to 4−hydroxy−L−proline
RNA splicing
SRP−dependent cotranslational protein targeting to membrane
protein import into mitochondrial matrix
protein insertion into mitochondrial inner membrane
positive regulation of establishment of protein localization to telomere
cholesterol efflux
vacuolar transport
toxin transport
protein targeting to ER
intra−Golgi vesicle−mediated transport
regulation of exocytosis
clathrin−dependent endocytosis
ribosomal large subunit export from nucleus
retrograde vesicle−mediated transport, Golgi to endoplasmic reticulum
endoplasmic reticulum to Golgi vesicle−mediated transport
protein localization to endosome
regulation of protein localization to late endosome
proton transmembrane transport
neurotransmitter receptor localization to postsynaptic specialization membrane
positive regulation of telomerase RNA localization to Cajal body
protein localization involved in protein targeting to mitochondrion
neurotransmitter receptor transport to postsynaptic membrane
vesicle−mediated transport in synapse
phospholipid efflux
intracellular protein transport
vesicle−mediated transport
amino acid transport
reverse cholesterol transport
vesicle docking involved in exocytosis
negative regulation of sodium ion transport
retrograde transport, endosome to Golgi
ribosomal large subunit assembly
mitochondrial respiratory chain complex I assembly
ribosomal small subunit assembly
extracellular matrix organization
proteasome assembly
rRNA processing
maturation of LSU−rRNA from tricistronic rRNA transcript (SSU−rRNA, 5.8S rRNA, LSU−rRNA)
membrane organization
mature ribosome assembly
platelet dense granule organization
Golgi organization
positive regulation of lipoprotein particle assembly
ribosomal large subunit biogenesis
negative regulation of epigenetic
maturation of SSU−rRNA from tricistronic rRNA transcript (SSU−rRNA, 5.8S rRNA, LSU−rRNA)
high−density lipoprotein particle assembly
cell junction assembly
regulation of protein−containing complex assembly
negative regulation of protein−containing complex assembly
eye pigment granule organization
negative regulation of focal adhesion assembly
collagen fibril organization
positive regulation of lamellipodium assembly
negative regulation of cell growth
positive regulation of extracellular matrix assembly
regulation of transcription from RNA polymerase II promoter in response to hypoxia
endoplasmic reticulum unfolded protein response
response to endoplasmic reticulum stress
cellular response to unfolded protein
cellular response to glucose starvation
regulation of transforming growth factor beta receptor signaling
IRE1−mediated unfolded protein response
response to starvation
cellular response to insulin stimulus
cellular response to BMP stimulus
cellular response to hypoxia
positive regulation of transforming growth factor beta receptor signaling pathway
cellular response to interleukin−7
negative regulation of IRE1−mediated unfolded protein response
cellular response to antibiotic
response to unfolded protein
negative regulation of endoplasmic reticulum unfolded protein response
response to nutrient
nuclear−transcribed mRNA catabolic process, nonsense−mediated decay
regulation of mRNA stability
autophagosome maturation
anaphase−promoting complex−dependent catabolic process
proteasomal ubiquitin−dependent protein catabolic process
proteolysis involved in protein catabolic process
late endosomal microautophagy
regulation of protein catabolic process
positive regulation of autophagy
proteasomal protein catabolic process
SCF−dependent proteasomal ubiquitin−dependent protein catabolic process
mRNA catabolic process
ubiquitin−dependent protein catabolic process
regulation of hematopoietic stem cell differentiation
decidualization
epithelial tube branching involved in lung morphogenesis
limb bud formation
multicellular organism aging
positive regulation of epithelial to mesenchymal transition
bone mineralization
positive regulation of fat cell differentiation
gland development
cell differentiation
neuron development
epidermis development
mesoderm formation
regulation of cell shape
positive regulation of dendritic spine development
striated muscle cell differentiation
transdifferentiation
NIK/NF−kappaB signaling
positive regulation of canonical Wnt signaling pathway
Rab protein signal transduction
integrin−mediated signaling pathway
positive regulation of signal transduction by p53 class mediator
phosphatidylinositol 3−kinase signaling
negative regulation of MAPK cascade
nodal signaling pathway
Wnt signaling pathway, planar cell polarity pathway
regulation of long−term synaptic potentiation
cell redox homeostasis
protein stabilization
cellular iron ion homeostasis
fatty acid homeostasis
cellular response to glucose stimulus
regulation of peroxisome size
antigen processing and presentation of exogenous peptide antigen via MHC class I, TAP−dependent
Fc−epsilon receptor signaling pathway
stimulatory C−type lectin receptor signaling pathway
natural killer cell mediated cytotoxicity
regulation of cellular amino acid metabolic process
D−xylose metabolic process
glycine biosynthetic process, by transamination of glyoxylate
fatty acid beta−oxidation using acyl−CoA dehydrogenase
citrate metabolic process
glycerol ether metabolic process
carbohydrate metabolic process
protein folding
protein refolding
'de novo' post−translational protein folding
chaperone−mediated protein folding
negative regulation of G2/M transition of mitotic cell cycle
positive regulation of epithelial cell migration
regulation of transforming growth factor beta2 production
viral transcription
positive regulation of epithelial cell proliferation
cell population proliferation
vesicle transport along microtubule
actin filament−based process