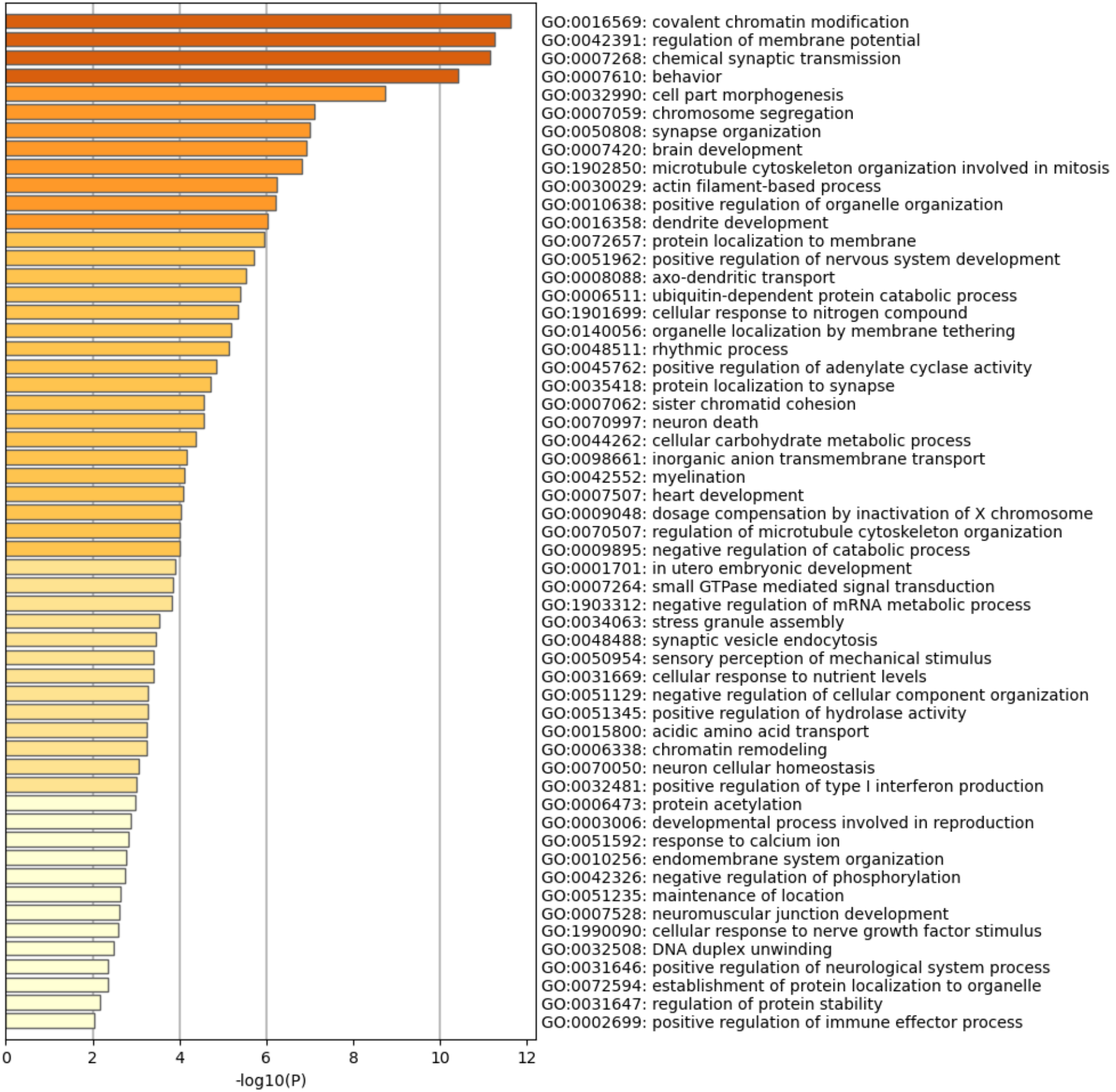
GO:0016569: covalent chromatin modification
GO:0042391: regulation of membrane potential
GO:0007268: chemical synaptic transmission
GO:0007610: behavior
GO:0032990: cell part morphogenesis
GO:0007059: chromosome segregation
GO:0050808: synapse organization
GO:0007420: brain development
GO:1902850: microtubule cytoskeleton organization involved in mitosis
GO:0030029: actin filament-based process
GO:0010638: positive regulation of organelle organization
GO:0016358: dendrite development
GO:0072657: protein localization to membrane
GO:0051962: positive regulation of nervous system development
GO:0008088: axo-dendritic transport
GO:0006511: ubiquitin-dependent protein catabolic process
GO:1901699: cellular response to nitrogen compound
GO:0140056: organelle localization by membrane tethering
GO:0048511: rhythmic process
GO:0045762: positive regulation of adenylate cyclase activity
GO:0035418: protein localization to synapse
GO:0007062: sister chromatid cohesion
GO:0070997: neuron death
GO:0044262: cellular carbohydrate metabolic process
GO:0098661: inorganic anion transmembrane transport
GO:0042552: myelination
GO:0007507: heart development
GO:0009048: dosage compensation by inactivation of X chromosome
GO:0070507: regulation of microtubule cytoskeleton organization
GO:0009895: negative regulation of catabolic process
GO:0001701: in utero embryonic development
GO:0007264: small GTPase mediated signal transduction
GO:1903312: negative regulation of mRNA metabolic process
GO:0034063: stress granule assembly
GO:0048488: synaptic vesicle endocytosis
GO:0050954: sensory perception of mechanical stimulus
GO:0031669: cellular response to nutrient levels
GO:0051129: negative regulation of cellular component organization
GO:0051345: positive regulation of hydrolase activity
GO:0015800: acidic amino acid transport
GO:0006338: chromatin remodeling
GO:0070050: neuron cellular homeostasis
GO:0032481: positive regulation of type I interferon production
GO:0006473: protein acetylation
GO:0003006: developmental process involved in reproduction
GO:0051592: response to calcium ion
GO:0010256: endomembrane system organization
GO:0042326: negative regulation of phosphorylation
GO:0051235: maintenance of location
GO:0007528: neuromuscular junction development
GO:1990090: cellular response to nerve growth factor stimulus
GO:0032508: DNA duplex unwinding
GO:0031646: positive regulation of neurological system process
GO:0072594: establishment of protein localization to organelle
GO:0031647: regulation of protein stability
GO:0002699: positive regulation of immune effector process
0
2
4
6
8
10
12
-log10(P)