

term	p-value	q-value
Thyroid Hormone Metabolic Process (GO:0042403)	0,000312	0,051678
Protein Localization to Endoplasmic Reticulum (GO:0070972)	0,000444	0,051678
Respiratory System Development (GO:0060541)	0,00127	0,0986
Regulation of Sprouting Angiogenesis (GO:1903670)	0,001788	0,104167
Antibacterial Humoral Response (GO:0019731)	0,003452	0,12699
Hydrogen Peroxide Biosynthetic Process (GO:0050665)	0,008224	0,12699
Intracellular Phosphate Ion Homeostasis (GO:0030643)	0,008224	0,12699
Sphingomyelin Biosynthetic Process (GO:0006686)	0,008224	0,12699
Protein Localization to Endoplasmic Reticulum Exit Site (GO:0070973)	0,011495	0,12699
Sodium-Dependent Phosphate Transport (GO:0044341)	0,011495	0,12699
Regulation of Protein Localization to Nucleolus (GO:1904749)	0,011495	0,12699
Reactive Oxygen Species Biosynthetic Process (GO:1903409)	0,013126	0,12699
Positive Regulation by Host of Viral Genome Replication (GO:0044829)	0,013126	0,12699
Endoplasmic Reticulum to Golgi Vesicle-Mediated Transport (GO:0006888)	0,014655	0,12699
Glucosamine-Containing Compound Catabolic Process (GO:1901072)	0,014755	0,12699
Granulocyte Migration (GO:0097530)	0,014755	0,12699
Activation of NF-kappaB-inducing Kinase Activity (GO:0007250)	0,014755	0,12699
Thyroid Hormone Generation (GO:0006590)	0,016382	0,12699
Glutamine Family Amino Acid Biosynthetic Process (GO:0009084)	0,016382	0,12699
Negative Regulation of Smooth Muscle Cell Apoptotic Process (GO:0034392)	0,016382	0,12699
Negative Regulation by Host of Viral Genome Replication (GO:0044828)	0,016382	0,12699
Negative Regulation of Vascular Endothelial Growth Factor Receptor Signaling Pathway (GO:0044828)	0,016382	0,12699
Positive Regulation of Defense Response to Bacterium (GO:1900426)	0,016382	0,12699
Positive Regulation of Myeloid Leukocyte Mediated Immunity (GO:0002888)	0,018005	0,12699
Negative Regulation of Sprouting Angiogenesis (GO:1903671)	0,018005	0,12699
Sphingomyelin Metabolic Process (GO:0006684)	0,019626	0,12699
Positive Regulation of Peptidyl-Threonine Phosphorylation (GO:0010800)	0,019626	0,12699
Granulocyte Differentiation (GO:0030851)	0,019626	0,12699
Phosphate Ion Homeostasis (GO:0055062)	0,019626	0,12699
Positive Regulation of Leukocyte Degranulation (GO:0043302)	0,019626	0,12699
Glutamine Metabolic Process (GO:0006541)	0,019626	0,12699
Plasma Membrane Bounded Cell Projection Organization (GO:0120036)	0,022591	0,12699
Negative Regulation of Muscle Cell Apoptotic Process (GO:0010656)	0,022861	0,12699
Glutamine Family Amino Acid Catabolic Process (GO:0009065)	0,022861	0,12699
Negative Regulation by Host of Viral Process (GO:0044793)	0,022861	0,12699
Synapse Organization (GO:0050808)	0,023494	0,12699
Antimicrobial Humoral Response (GO:0019730)	0,024411	0,12699
Ceramide Transport (GO:0035627)	0,024474	0,12699
Dicarboxylic Acid Catabolic Process (GO:0043649)	0,024474	0,12699
Regulation of Smooth Muscle Cell Apoptotic Process (GO:0034391)	0,026085	0,12699
Regulation of Superoxide Anion Generation (GO:0032928)	0,026085	0,12699
Positive Regulation of ERAD Pathway (GO:1904294)	0,026085	0,12699
Positive Regulation of Superoxide Anion Generation (GO:0032930)	0,026085	0,12699
Cell Junction Maintenance (GO:0034331)	0,027694	0,12699
Protein Autoprocessing (GO:0016540)	0,027694	0,12699
Neuron Development (GO:0048666)	0,028227	0,12699
Phosphate Ion Transport (GO:0006817)	0,029299	0,12699
Positive Regulation by Host of Viral Process (GO:0044794)	0,029299	0,12699
Amyloid Precursor Protein Metabolic Process (GO:0042982)	0,029299	0,12699
Maintenance of Gastrointestinal Epithelium (GO:0030277)	0,030902	0,12699
Insulin Secretion (GO:0030073)	0,030902	0,12699
Regulation of Peptidyl-Threonine Phosphorylation (GO:0010799)	0,030902	0,12699

Superoxide Anion Generation (GO:0042554)	0,030902	0,12699
Cell-Cell Adhesion via Plasma-Membrane Adhesion Molecules (GO:0098742)	0,032266	0,12699
Glutamate Metabolic Process (GO:0006536)	0,032503	0,12699
Peptide Cross-Linking (GO:0018149)	0,034101	0,12699
Epithelial Structure Maintenance (GO:0010669)	0,035696	0,12699
Alpha-Amino Acid Catabolic Process (GO:1901606)	0,035696	0,12699
Regulation of Neurotransmitter Receptor Activity (GO:0099601)	0,037289	0,12699
Modulation by Host of Viral Genome Replication (GO:0044827)	0,037289	0,12699
Protein Palmitoylation (GO:0018345)	0,037289	0,12699
Regulation of ERAD Pathway (GO:1904292)	0,037289	0,12699
Protein Localization to Cell Surface (GO:0034394)	0,038879	0,12699
Cellular Response to Acetylcholine (GO:1905145)	0,038879	0,12699
Secondary Alcohol Biosynthetic Process (GO:1902653)	0,038879	0,12699
Organophosphate Ester Transport (GO:0015748)	0,040467	0,12699
Protein Lipidation (GO:0006497)	0,040467	0,12699
Cholesterol Biosynthetic Process (GO:0006695)	0,040467	0,12699
Response to Purine-Containing Compound (GO:0014074)	0,040467	0,12699
Non-Canonical NF-kappaB Signal Transduction (GO:0038061)	0,040467	0,12699
Regulation of Vascular Endothelial Growth Factor Receptor Signaling Pathway (GO:0030947)	0,042052	0,12699
Respiratory Tube Development (GO:0030323)	0,042052	0,12699
Sterol Transport (GO:0015918)	0,042052	0,12699
Positive Regulation of Erythrocyte Differentiation (GO:0045648)	0,043635	0,12699
Amide Transport (GO:0042886)	0,043635	0,12699
Peptide Hormone Secretion (GO:0030072)	0,043635	0,12699
Membrane Fusion (GO:0061025)	0,045215	0,12699
Phospholipid Biosynthetic Process (GO:0008654)	0,045215	0,12699
Positive Regulation of G Protein-Coupled Receptor Signaling Pathway (GO:0045745)	0,045215	0,12699
Hydrogen Peroxide Metabolic Process (GO:0042743)	0,045215	0,12699
Positive Regulation of Response to Endoplasmic Reticulum Stress (GO:1905898)	0,045215	0,12699
Sterol Biosynthetic Process (GO:0016126)	0,045215	0,12699
Neuron Projection Development (GO:0031175)	0,045237	0,12699
Regulation of Signal Transduction (GO:0009966)	0,047225	0,129452
Regulation of Angiogenesis (GO:0045765)	0,047225	0,129452
Positive Regulation of Response to Wounding (GO:1903036)	0,048368	0,131043
Collagen Catabolic Process (GO:0030574)	0,04994	0,131889
Leukocyte Cell-Cell Adhesion (GO:0007159)	0,04994	0,131889