**Supplementary Table 7.** **Top gene sets and pathways enriched in PASCAL.**

|  |  |
| --- | --- |
| Pathway ID | P |
| GO\_ANATOMICAL\_STRUCTURE\_FORMATION\_INVOLVED\_IN\_MORPHOGENESIS | < 1.00E-16 |
| GO\_ION\_TRANSPORT | < 1.00E-16 |
| GO\_NEUROLOGICAL\_SYSTEM\_PROCESS | < 1.00E-16 |
| GO\_NEGATIVE\_REGULATION\_OF\_GENE\_EXPRESSION | < 1.00E-16 |
| GO\_CHEMICAL\_HOMEOSTASIS | < 1.00E-16 |
| GO\_IMMUNE\_SYSTEM\_PROCESS | < 1.00E-16 |
| GO\_POSITIVE\_REGULATION\_OF\_RESPONSE\_TO\_STIMULUS | < 1.00E-16 |
| GO\_POSITIVE\_REGULATION\_OF\_CELL\_COMMUNICATION | < 1.00E-16 |
| GO\_REGULATION\_OF\_HYDROLASE\_ACTIVITY | < 1.00E-16 |
| GO\_CELL\_CELL\_SIGNALING | < 1.00E-16 |
| GO\_SMALL\_MOLECULE\_METABOLIC\_PROCESS | < 1.00E-16 |
| GO\_HOMEOSTATIC\_PROCESS | < 1.00E-16 |
| GO\_REGULATION\_OF\_INTRACELLULAR\_SIGNAL\_TRANSDUCTION | < 1.00E-16 |
| GO\_POSITIVE\_REGULATION\_OF\_BIOSYNTHETIC\_PROCESS | < 1.00E-16 |
| GO\_NEGATIVE\_REGULATION\_OF\_NITROGEN\_COMPOUND\_METABOLIC\_PROCESS | < 1.00E-16 |
| GO\_REGULATION\_OF\_NERVOUS\_SYSTEM\_DEVELOPMENT | < 1.00E-16 |
| GO\_ESTABLISHMENT\_OF\_LOCALIZATION\_IN\_CELL | < 1.00E-16 |
| GO\_VESICLE\_MEDIATED\_TRANSPORT | < 1.00E-16 |
| GO\_POSITIVE\_REGULATION\_OF\_CELLULAR\_COMPONENT\_ORGANIZATION | < 1.00E-16 |
| GO\_POSITIVE\_REGULATION\_OF\_GENE\_EXPRESSION | < 1.00E-16 |
| GO\_CARBOHYDRATE\_DERIVATIVE\_METABOLIC\_PROCESS | < 1.00E-16 |
| GO\_CATION\_TRANSPORT | < 1.00E-16 |
| GO\_POSITIVE\_REGULATION\_OF\_HYDROLASE\_ACTIVITY | < 1.00E-16 |
| GO\_RESPONSE\_TO\_ENDOGENOUS\_STIMULUS | < 1.00E-16 |
| GO\_POSITIVE\_REGULATION\_OF\_CATALYTIC\_ACTIVITY | < 1.00E-16 |
| GO\_RESPONSE\_TO\_EXTERNAL\_STIMULUS | < 1.00E-16 |
| GO\_ION\_TRANSMEMBRANE\_TRANSPORT | < 1.00E-16 |
| GO\_INTRACELLULAR\_SIGNAL\_TRANSDUCTION | < 1.00E-16 |
| GO\_REGULATION\_OF\_CELL\_PROJECTION\_ORGANIZATION | < 1.00E-16 |
| GO\_POSITIVE\_REGULATION\_OF\_MOLECULAR\_FUNCTION | < 1.00E-16 |
| GO\_CELLULAR\_RESPONSE\_TO\_ORGANIC\_SUBSTANCE | < 1.00E-16 |
| GO\_POSITIVE\_REGULATION\_OF\_PHOSPHORUS\_METABOLIC\_PROCESS | < 1.00E-16 |
| GO\_REGULATION\_OF\_CELLULAR\_LOCALIZATION | < 1.00E-16 |
| GO\_REGULATION\_OF\_MULTICELLULAR\_ORGANISMAL\_DEVELOPMENT | < 1.00E-16 |
| GO\_MACROMOLECULAR\_COMPLEX\_ASSEMBLY | < 1.00E-16 |
| GO\_ORGAN\_MORPHOGENESIS | < 1.00E-16 |
| GO\_POSITIVE\_REGULATION\_OF\_MULTICELLULAR\_ORGANISMAL\_PROCESS | < 1.00E-16 |
| GO\_G\_PROTEIN\_COUPLED\_RECEPTOR\_SIGNALING\_PATHWAY | < 1.00E-16 |
| GO\_REGULATION\_OF\_RESPONSE\_TO\_STRESS | < 1.00E-16 |
| GO\_PHOSPHATE\_CONTAINING\_COMPOUND\_METABOLIC\_PROCESS | < 1.00E-16 |
| GO\_REGULATION\_OF\_CELL\_DIFFERENTIATION | < 1.00E-16 |
| GO\_REGULATION\_OF\_IMMUNE\_SYSTEM\_PROCESS | < 1.00E-16 |
| GO\_RESPONSE\_TO\_OXYGEN\_CONTAINING\_COMPOUND | < 1.00E-16 |
| GO\_LOCOMOTION | < 1.00E-16 |
| GO\_NEUROGENESIS | < 1.00E-16 |
| GO\_SYSTEM\_PROCESS | < 1.00E-16 |
| GO\_MOVEMENT\_OF\_CELL\_OR\_SUBCELLULAR\_COMPONENT | < 1.00E-16 |
| GO\_POSITIVE\_REGULATION\_OF\_PROTEIN\_METABOLIC\_PROCESS | < 1.00E-16 |
| GO\_RESPONSE\_TO\_ABIOTIC\_STIMULUS | < 1.00E-16 |
| GO\_POSITIVE\_REGULATION\_OF\_INTRACELLULAR\_SIGNAL\_TRANSDUCTION | < 1.00E-16 |
| GO\_HEAD\_DEVELOPMENT | < 1.00E-16 |
| GO\_ORGANONITROGEN\_COMPOUND\_METABOLIC\_PROCESS | < 1.00E-16 |
| GO\_POSITIVE\_REGULATION\_OF\_PROTEIN\_MODIFICATION\_PROCESS | < 1.00E-16 |
| GO\_NEGATIVE\_REGULATION\_OF\_DEVELOPMENTAL\_PROCESS | < 1.00E-16 |
| GO\_REGULATION\_OF\_ANATOMICAL\_STRUCTURE\_MORPHOGENESIS | < 1.00E-16 |
| GO\_REGULATION\_OF\_CELL\_DEATH | < 1.00E-16 |
| GO\_NEGATIVE\_REGULATION\_OF\_RESPONSE\_TO\_STIMULUS | < 1.00E-16 |
| GO\_PROTEIN\_LOCALIZATION | < 1.00E-16 |
| GO\_CELLULAR\_RESPONSE\_TO\_ENDOGENOUS\_STIMULUS | < 1.00E-16 |
| GO\_CELL\_PROJECTION\_ORGANIZATION | < 1.00E-16 |
| GO\_NEGATIVE\_REGULATION\_OF\_PROTEIN\_METABOLIC\_PROCESS | < 1.00E-16 |
| GO\_PROTEIN\_COMPLEX\_SUBUNIT\_ORGANIZATION | < 1.00E-16 |
| GO\_NEURON\_DIFFERENTIATION | < 1.00E-16 |
| GO\_POSITIVE\_REGULATION\_OF\_DEVELOPMENTAL\_PROCESS | < 1.00E-16 |
| GO\_REGULATION\_OF\_PROTEIN\_MODIFICATION\_PROCESS | < 1.00E-16 |
| GO\_POSITIVE\_REGULATION\_OF\_TRANSPORT | < 1.00E-16 |
| GO\_REGULATION\_OF\_TRANSPORT | < 1.00E-16 |
| GO\_POSITIVE\_REGULATION\_OF\_CELL\_DIFFERENTIATION | < 1.00E-16 |
| GO\_REGULATION\_OF\_ION\_TRANSPORT | < 1.00E-16 |
| GO\_CELLULAR\_RESPONSE\_TO\_STRESS | < 1.00E-16 |
| GO\_CELL\_DEVELOPMENT | < 1.00E-16 |
| GO\_REGULATION\_OF\_ORGANELLE\_ORGANIZATION | < 1.00E-16 |
| GO\_PHOSPHORYLATION | < 1.00E-16 |
| GO\_NEGATIVE\_REGULATION\_OF\_MULTICELLULAR\_ORGANISMAL\_PROCESS | < 1.00E-16 |
| GO\_REGULATION\_OF\_CELL\_DEVELOPMENT | < 1.00E-16 |
| GO\_CATABOLIC\_PROCESS | < 1.00E-16 |
| GO\_TRANSMEMBRANE\_TRANSPORT | < 1.00E-16 |
| GO\_INORGANIC\_ION\_TRANSMEMBRANE\_TRANSPORT | < 1.00E-16 |
| GO\_CENTRAL\_NERVOUS\_SYSTEM\_DEVELOPMENT | < 1.00E-16 |
| GO\_REGULATION\_OF\_PHOSPHORUS\_METABOLIC\_PROCESS | < 1.00E-16 |
| GO\_REGULATION\_OF\_CELL\_PROLIFERATION | < 1.00E-16 |
| GO\_BIOLOGICAL\_ADHESION | < 1.00E-16 |
| GO\_TISSUE\_DEVELOPMENT | < 1.00E-16 |
| GO\_NEGATIVE\_REGULATION\_OF\_CELL\_COMMUNICATION | < 1.00E-16 |
| GO\_REGULATION\_OF\_TRANSCRIPTION\_FROM\_RNA\_POLYMERASE\_II\_PROMOTER | < 1.00E-16 |
| GO\_VACUOLE | < 1.00E-16 |
| GO\_NEURON\_PART | < 1.00E-16 |
| GO\_INTRINSIC\_COMPONENT\_OF\_PLASMA\_MEMBRANE | < 1.00E-16 |
| GO\_CELL\_JUNCTION | < 1.00E-16 |
| GO\_CYTOSKELETON | < 1.00E-16 |
| GO\_INTRACELLULAR\_VESICLE | < 1.00E-16 |
| GO\_SYNAPSE | < 1.00E-16 |
| GO\_NEURON\_PROJECTION | < 1.00E-16 |
| GO\_EXTRACELLULAR\_SPACE | < 1.00E-16 |
| GO\_MITOCHONDRION | < 1.00E-16 |
| GO\_CELL\_SURFACE | < 1.00E-16 |
| GO\_TRANSPORTER\_COMPLEX | < 1.00E-16 |
| GO\_MEMBRANE\_PROTEIN\_COMPLEX | < 1.00E-16 |
| GO\_PLASMA\_MEMBRANE\_REGION | < 1.00E-16 |
| GO\_CELL\_PROJECTION | < 1.00E-16 |
| GO\_CELL\_PROJECTION\_PART | < 1.00E-16 |
| GO\_GOLGI\_APPARATUS | < 1.00E-16 |
| GO\_ENDOPLASMIC\_RETICULUM | < 1.00E-16 |
| GO\_MEMBRANE\_REGION | < 1.00E-16 |
| GO\_SYNAPSE\_PART | < 1.00E-16 |
| GO\_PLASMA\_MEMBRANE\_PROTEIN\_COMPLEX | < 1.00E-16 |
| GO\_SIGNAL\_TRANSDUCER\_ACTIVITY | < 1.00E-16 |
| GO\_CYTOSKELETAL\_PROTEIN\_BINDING | < 1.00E-16 |
| GO\_PROTEIN\_DIMERIZATION\_ACTIVITY | < 1.00E-16 |
| GO\_GATED\_CHANNEL\_ACTIVITY | < 1.00E-16 |
| GO\_PASSIVE\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY | < 1.00E-16 |
| GO\_CALCIUM\_ION\_BINDING | < 1.00E-16 |
| GO\_SIGNALING\_RECEPTOR\_ACTIVITY | < 1.00E-16 |
| GO\_NUCLEIC\_ACID\_BINDING\_TRANSCRIPTION\_FACTOR\_ACTIVITY | < 1.00E-16 |
| GO\_MOLECULAR\_FUNCTION\_REGULATOR | < 1.00E-16 |
| GO\_IDENTICAL\_PROTEIN\_BINDING | < 1.00E-16 |
| GO\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY | < 1.00E-16 |
| GO\_RIBONUCLEOTIDE\_BINDING | < 1.00E-16 |
| GO\_ENZYME\_REGULATOR\_ACTIVITY | < 1.00E-16 |
| GO\_ENZYME\_BINDING | < 1.00E-16 |
| GO\_RECEPTOR\_ACTIVITY | < 1.00E-16 |
| GO\_ADENYL\_NUCLEOTIDE\_BINDING | < 1.00E-16 |
| GO\_RECEPTOR\_BINDING | < 1.00E-16 |
| GO\_TRANSPORTER\_ACTIVITY | < 1.00E-16 |
| GO\_MACROMOLECULAR\_COMPLEX\_BINDING | < 1.00E-16 |
| GO\_NEURON\_DEVELOPMENT | 1.11E-16 |
| GO\_CELL\_DEATH | 1.11E-16 |
| GO\_POSITIVE\_REGULATION\_OF\_CELL\_PROLIFERATION | 1.11E-16 |
| GO\_REGULATION\_OF\_NEURON\_DIFFERENTIATION | 1.11E-16 |
| GO\_REGULATION\_OF\_GTPASE\_ACTIVITY | 1.11E-16 |
| GO\_SOMATODENDRITIC\_COMPARTMENT | 1.11E-16 |
| GO\_SEQUENCE\_SPECIFIC\_DNA\_BINDING | 1.11E-16 |
| GO\_TRANSFERASE\_ACTIVITY\_TRANSFERRING\_PHOSPHORUS\_CONTAINING\_GROUPS | 1.11E-16 |
| GO\_EPITHELIUM\_DEVELOPMENT | 2.22E-16 |
| GO\_MEMBRANE\_ORGANIZATION | 2.22E-16 |
| GO\_GOLGI\_APPARATUS\_PART | 2.22E-16 |
| GO\_NEGATIVE\_REGULATION\_OF\_MOLECULAR\_FUNCTION | 5.55E-16 |
| GO\_LIPID\_METABOLIC\_PROCESS | 5.55E-16 |
| GO\_ESTABLISHMENT\_OF\_PROTEIN\_LOCALIZATION | 8.88E-16 |
| GO\_TRANSITION\_METAL\_ION\_BINDING | 1.44E-15 |
| GO\_METAL\_ION\_TRANSPORT | 1.78E-15 |
| GO\_PROTEIN\_COMPLEX\_BIOGENESIS | 2.11E-15 |
| GO\_CELL\_MOTILITY | 3.55E-15 |
| GO\_ENDOPLASMIC\_RETICULUM\_PART | 3.66E-15 |
| GO\_REGULATION\_OF\_CELLULAR\_COMPONENT\_MOVEMENT | 6.44E-15 |
| GO\_REGULATION\_OF\_RESPONSE\_TO\_EXTERNAL\_STIMULUS | 7.11E-15 |
| GO\_PROTEIN\_COMPLEX\_BINDING | 8.22E-15 |
| GO\_METAL\_ION\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY | 8.33E-15 |
| GO\_CELLULAR\_COMPONENT\_MORPHOGENESIS | 8.77E-15 |
| GO\_CELL\_CELL\_ADHESION | 1.34E-14 |
| GO\_CATION\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY | 1.34E-14 |
| GO\_KINASE\_ACTIVITY | 1.92E-14 |
| GO\_CELLULAR\_HOMEOSTASIS | 2.29E-14 |
| GO\_NEGATIVE\_REGULATION\_OF\_CELL\_DEATH | 2.94E-14 |
| GO\_CELLULAR\_MACROMOLECULE\_LOCALIZATION | 3.46E-14 |
| GO\_CYTOSKELETAL\_PART | 3.69E-14 |
| GO\_REGULATION\_OF\_TRANSFERASE\_ACTIVITY | 6.03E-14 |
| GO\_REGULATION\_OF\_TRANSMEMBRANE\_TRANSPORT | 7.44E-14 |
| GO\_REGULATION\_OF\_NEURON\_PROJECTION\_DEVELOPMENT | 8.23E-14 |
| GO\_REGULATION\_OF\_CELL\_MORPHOGENESIS | 8.47E-14 |
| GO\_ENZYME\_LINKED\_RECEPTOR\_PROTEIN\_SIGNALING\_PATHWAY | 1.09E-13 |
| GO\_INORGANIC\_CATION\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY | 1.42E-13 |
| REACTOME\_SIGNALING\_BY\_GPCR | 1.48E-13 |
| GO\_CATION\_CHANNEL\_ACTIVITY | 1.83E-13 |
| GO\_SENSORY\_PERCEPTION | 1.87E-13 |
| GO\_REPRODUCTION | 1.92E-13 |
| GO\_IMMUNE\_RESPONSE | 2.09E-13 |
| GO\_POSITIVE\_REGULATION\_OF\_TRANSCRIPTION\_FROM\_RNA\_POLYMERASE\_II\_PROMOTER | 2.12E-13 |
| GO\_SYNAPTIC\_SIGNALING | 2.30E-13 |
| GO\_RNA\_BINDING | 2.33E-13 |
| GO\_PROTEOLYSIS | 4.98E-13 |
| REACTOME\_GPCR\_DOWNSTREAM\_SIGNALING | 6.28E-13 |
| GO\_CATION\_CHANNEL\_COMPLEX | 7.46E-13 |
| GO\_POSITIVE\_REGULATION\_OF\_IMMUNE\_SYSTEM\_PROCESS | 1.51E-12 |
| GO\_SINGLE\_ORGANISM\_CELLULAR\_LOCALIZATION | 1.61E-12 |
| GO\_BEHAVIOR | 1.92E-12 |
| GO\_REGULATION\_OF\_CELLULAR\_COMPONENT\_BIOGENESIS | 2.22E-12 |
| GO\_AXON | 3.06E-12 |
| GO\_RESPONSE\_TO\_ORGANIC\_CYCLIC\_COMPOUND | 3.11E-12 |
| GO\_DEFENSE\_RESPONSE | 3.14E-12 |
| GO\_CIRCULATORY\_SYSTEM\_DEVELOPMENT | 3.31E-12 |
| GO\_REGULATION\_OF\_CELL\_ADHESION | 3.60E-12 |
| GO\_NEGATIVE\_REGULATION\_OF\_CELL\_DIFFERENTIATION | 3.74E-12 |
| GO\_SECRETION | 4.56E-12 |
| GO\_PROTEIN\_PHOSPHORYLATION | 5.36E-12 |
| GO\_NEURON\_PROJECTION\_DEVELOPMENT | 5.65E-12 |
| GO\_PERINUCLEAR\_REGION\_OF\_CYTOPLASM | 5.69E-12 |
| GO\_REGULATION\_OF\_SECRETION | 5.79E-12 |
| GO\_CELLULAR\_RESPONSE\_TO\_OXYGEN\_CONTAINING\_COMPOUND | 6.91E-12 |
| GO\_REGULATION\_OF\_PROTEIN\_LOCALIZATION | 7.59E-12 |
| GO\_DENDRITE | 1.08E-11 |
| GO\_SINGLE\_ORGANISM\_BIOSYNTHETIC\_PROCESS | 1.09E-11 |
| GO\_REGULATION\_OF\_SYSTEM\_PROCESS | 1.33E-11 |
| GO\_REGULATORY\_REGION\_NUCLEIC\_ACID\_BINDING | 1.56E-11 |
| GO\_POSITIVE\_REGULATION\_OF\_CELL\_DEVELOPMENT | 1.59E-11 |
| GO\_RECEPTOR\_COMPLEX | 1.80E-11 |
| GO\_POSITIVE\_REGULATION\_OF\_NERVOUS\_SYSTEM\_DEVELOPMENT | 2.38E-11 |
| GO\_REGULATION\_OF\_KINASE\_ACTIVITY | 2.53E-11 |
| GO\_CELL\_MORPHOGENESIS\_INVOLVED\_IN\_DIFFERENTIATION | 2.60E-11 |
| GO\_SENSORY\_ORGAN\_DEVELOPMENT | 2.76E-11 |
| GO\_POSTSYNAPSE | 3.21E-11 |
| GO\_REGULATION\_OF\_IMMUNE\_RESPONSE | 3.66E-11 |
| GO\_CALCIUM\_ION\_TRANSPORT | 4.61E-11 |
| GO\_POSITIVE\_REGULATION\_OF\_CELL\_PROJECTION\_ORGANIZATION | 4.66E-11 |
| GO\_DIVALENT\_INORGANIC\_CATION\_TRANSPORT | 5.24E-11 |
| GO\_CELLULAR\_CHEMICAL\_HOMEOSTASIS | 7.68E-11 |
| GO\_ANCHORING\_JUNCTION | 8.70E-11 |
| GO\_ORGANOPHOSPHATE\_METABOLIC\_PROCESS | 9.15E-11 |
| GO\_RESPONSE\_TO\_NITROGEN\_COMPOUND | 1.11E-10 |
| GO\_CELL\_ACTIVATION | 1.24E-10 |
| GO\_MULTICELLULAR\_ORGANISMAL\_SIGNALING | 1.28E-10 |
| GO\_LIGAND\_GATED\_CHANNEL\_ACTIVITY | 1.33E-10 |
| GO\_REGULATION\_OF\_MEMBRANE\_POTENTIAL | 1.34E-10 |
| GO\_NEGATIVE\_REGULATION\_OF\_CATALYTIC\_ACTIVITY | 1.39E-10 |
| GO\_CELLULAR\_CATABOLIC\_PROCESS | 1.41E-10 |
| GO\_REGULATION\_OF\_METAL\_ION\_TRANSPORT | 1.49E-10 |
| REACTOME\_NEURONAL\_SYSTEM | 2.04E-10 |
| GO\_SECRETION\_BY\_CELL | 2.15E-10 |
| GO\_NEGATIVE\_REGULATION\_OF\_TRANSCRIPTION\_FROM\_RNA\_POLYMERASE\_II\_PROMOTER | 2.43E-10 |
| GO\_CELL\_PART\_MORPHOGENESIS | 2.63E-10 |
| GO\_SINGLE\_ORGANISM\_CATABOLIC\_PROCESS | 2.76E-10 |
| GO\_SYNAPTIC\_MEMBRANE | 2.92E-10 |
| GO\_PROTEINACEOUS\_EXTRACELLULAR\_MATRIX | 2.99E-10 |
| GO\_GOLGI\_MEMBRANE | 3.34E-10 |
| GO\_EXTRACELLULAR\_MATRIX | 3.53E-10 |
| GO\_CALCIUM\_ION\_TRANSMEMBRANE\_TRANSPORT | 3.74E-10 |
| GO\_POSITIVE\_REGULATION\_OF\_NEURON\_DIFFERENTIATION | 4.59E-10 |
| GO\_POSITIVE\_REGULATION\_OF\_NEURON\_PROJECTION\_DEVELOPMENT | 4.84E-10 |
| GO\_EMBRYO\_DEVELOPMENT | 5.43E-10 |
| GO\_IMMUNE\_SYSTEM\_DEVELOPMENT | 5.50E-10 |
| GO\_REGULATION\_OF\_HEART\_CONTRACTION | 6.02E-10 |
| GO\_REGULATION\_OF\_BLOOD\_CIRCULATION | 6.09E-10 |
| GO\_OXIDATION\_REDUCTION\_PROCESS | 6.22E-10 |
| GO\_CELL\_MORPHOGENESIS\_INVOLVED\_IN\_NEURON\_DIFFERENTIATION | 7.24E-10 |
| GO\_ZINC\_ION\_BINDING | 7.50E-10 |
| GO\_REGULATION\_OF\_CELL\_CYCLE | 8.19E-10 |
| GO\_CALCIUM\_ION\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY | 8.92E-10 |
| GO\_PROTEIN\_DOMAIN\_SPECIFIC\_BINDING | 9.15E-10 |
| GO\_REGULATION\_OF\_VESICLE\_MEDIATED\_TRANSPORT | 9.68E-10 |
| GO\_SENSORY\_PERCEPTION\_OF\_MECHANICAL\_STIMULUS | 1.06E-09 |
| GO\_CELL\_CYCLE | 1.31E-09 |
| GO\_CYTOSKELETON\_ORGANIZATION | 1.34E-09 |
| GO\_DIVALENT\_INORGANIC\_CATION\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY | 1.40E-09 |
| GO\_DOUBLE\_STRANDED\_DNA\_BINDING | 2.38E-09 |
| GO\_ENDOSOME | 2.42E-09 |
| GO\_POSITIVE\_REGULATION\_OF\_TRANSFERASE\_ACTIVITY | 2.43E-09 |
| GO\_NEURON\_PROJECTION\_MORPHOGENESIS | 3.08E-09 |
| GO\_CELL\_BODY | 3.39E-09 |
| GO\_REGULATION\_OF\_GROWTH | 3.46E-09 |
| KEGG\_NEUROACTIVE\_LIGAND\_RECEPTOR\_INTERACTION | 3.67E-09 |
| GO\_POTASSIUM\_CHANNEL\_COMPLEX | 4.35E-09 |
| GO\_CIRCULATORY\_SYSTEM\_PROCESS | 5.69E-09 |
| GO\_CELLULAR\_LIPID\_METABOLIC\_PROCESS | 5.81E-09 |
| GO\_HYDROLASE\_ACTIVITY\_ACTING\_ON\_ESTER\_BONDS | 5.86E-09 |
| GO\_PROTEIN\_HOMODIMERIZATION\_ACTIVITY | 6.74E-09 |
| GO\_CELL\_CELL\_ADHESION\_VIA\_PLASMA\_MEMBRANE\_ADHESION\_MOLECULES | 7.69E-09 |
| GO\_FOREBRAIN\_DEVELOPMENT | 8.22E-09 |
| GO\_G\_PROTEIN\_COUPLED\_RECEPTOR\_ACTIVITY | 8.82E-09 |
| GO\_ORGANIC\_ACID\_METABOLIC\_PROCESS | 9.62E-09 |
| GO\_ENZYME\_ACTIVATOR\_ACTIVITY | 1.02E-08 |
| GO\_ACTIN\_BINDING | 1.05E-08 |
| GO\_TAXIS | 1.14E-08 |
| REACTOME\_DEVELOPMENTAL\_BIOLOGY | 1.15E-08 |
| GO\_ION\_HOMEOSTASIS | 1.16E-08 |
| GO\_RESPONSE\_TO\_WOUNDING | 1.18E-08 |
| GO\_REGULATION\_OF\_BODY\_FLUID\_LEVELS | 1.48E-08 |
| GO\_POLY\_A\_RNA\_BINDING | 1.51E-08 |
| GO\_REGULATION\_OF\_ACTIN\_FILAMENT\_BASED\_PROCESS | 1.56E-08 |
| GO\_OXIDOREDUCTASE\_ACTIVITY | 1.59E-08 |
| GO\_CATALYTIC\_COMPLEX | 2.01E-08 |
| GO\_NUCLEAR\_OUTER\_MEMBRANE\_ENDOPLASMIC\_RETICULUM\_MEMBRANE\_NETWORK | 2.07E-08 |
| GO\_HOMOPHILIC\_CELL\_ADHESION\_VIA\_PLASMA\_MEMBRANE\_ADHESION\_MOLECULES | 2.12E-08 |
| GO\_SKELETAL\_SYSTEM\_DEVELOPMENT | 2.13E-08 |
| GO\_REGULATION\_OF\_CELL\_MORPHOGENESIS\_INVOLVED\_IN\_DIFFERENTIATION | 2.20E-08 |
| GO\_REGULATION\_OF\_MAPK\_CASCADE | 2.31E-08 |
| GO\_TRANSMEMBRANE\_RECEPTOR\_PROTEIN\_TYROSINE\_KINASE\_SIGNALING\_PATHWAY | 2.63E-08 |
| GO\_POSTSYNAPTIC\_MEMBRANE | 2.79E-08 |
| GO\_RESPONSE\_TO\_HORMONE | 2.91E-08 |
| GO\_REGULATION\_OF\_ANATOMICAL\_STRUCTURE\_SIZE | 2.97E-08 |
| GO\_PRESYNAPSE | 3.06E-08 |
| GO\_MICROTUBULE\_CYTOSKELETON | 3.72E-08 |
| GO\_MUSCLE\_SYSTEM\_PROCESS | 3.74E-08 |
| GO\_TELENCEPHALON\_DEVELOPMENT | 4.12E-08 |
| REACTOME\_TRANSMEMBRANE\_TRANSPORT\_OF\_SMALL\_MOLECULES | 4.46E-08 |
| GO\_PROTEIN\_KINASE\_ACTIVITY | 5.03E-08 |
| GO\_LOCOMOTORY\_BEHAVIOR | 5.36E-08 |
| GO\_ACTIN\_FILAMENT\_BASED\_PROCESS | 5.86E-08 |
| GO\_CARDIAC\_CONDUCTION | 6.05E-08 |
| GO\_REGULATION\_OF\_CATION\_TRANSMEMBRANE\_TRANSPORT | 6.06E-08 |
| GO\_SENSORY\_PERCEPTION\_OF\_CHEMICAL\_STIMULUS | 6.41E-08 |
| GO\_NEGATIVE\_REGULATION\_OF\_CELLULAR\_COMPONENT\_ORGANIZATION | 6.63E-08 |
| GO\_POSITIVE\_REGULATION\_OF\_KINASE\_ACTIVITY | 7.33E-08 |
| GO\_RNA\_POLYMERASE\_II\_TRANSCRIPTION\_FACTOR\_ACTIVITY\_SEQUENCE\_SPECIFIC\_DNA\_BINDING | 8.93E-08 |
| GO\_ANION\_TRANSPORT | 9.19E-08 |
| GO\_ENDOCYTOSIS | 9.78E-08 |
| GO\_ACTIN\_CYTOSKELETON | 1.04E-07 |
| GO\_HYDROLASE\_ACTIVITY\_ACTING\_ON\_ACID\_ANHYDRIDES | 1.09E-07 |
| GO\_NUCLEOSIDE\_TRIPHOSPHATASE\_REGULATOR\_ACTIVITY | 1.14E-07 |
| GO\_MUSCLE\_CONTRACTION | 1.41E-07 |
| GO\_APICAL\_PART\_OF\_CELL | 1.43E-07 |
| GO\_MODULATION\_OF\_SYNAPTIC\_TRANSMISSION | 1.44E-07 |
| GO\_KINASE\_BINDING | 1.63E-07 |
| GO\_WOUND\_HEALING | 1.81E-07 |
| GO\_HEMOSTASIS | 1.91E-07 |
| GO\_NEURON\_RECOGNITION | 2.10E-07 |
| GO\_REGULATION\_OF\_DEFENSE\_RESPONSE | 2.12E-07 |
| REACTOME\_ION\_CHANNEL\_TRANSPORT | 2.16E-07 |
| GO\_GUANYL\_NUCLEOTIDE\_EXCHANGE\_FACTOR\_ACTIVITY | 2.20E-07 |
| GO\_NEURON\_PROJECTION\_GUIDANCE | 2.40E-07 |
| GO\_SIDE\_OF\_MEMBRANE | 2.70E-07 |
| GO\_RESPONSE\_TO\_CYTOKINE | 2.75E-07 |
| GO\_CELLULAR\_RESPONSE\_TO\_NITROGEN\_COMPOUND | 2.87E-07 |
| GO\_SINGLE\_ORGANISM\_BEHAVIOR | 2.91E-07 |
| GO\_ORGANONITROGEN\_COMPOUND\_BIOSYNTHETIC\_PROCESS | 2.93E-07 |
| GO\_SINGLE\_ORGANISM\_CELL\_ADHESION | 2.97E-07 |
| GO\_ANCHORED\_COMPONENT\_OF\_MEMBRANE | 2.99E-07 |
| GO\_TRANSCRIPTION\_FROM\_RNA\_POLYMERASE\_II\_PROMOTER | 3.19E-07 |
| GO\_CELL\_LEADING\_EDGE | 3.40E-07 |
| GO\_REGULATION\_OF\_HORMONE\_LEVELS | 3.56E-07 |
| GO\_NEGATIVE\_REGULATION\_OF\_PROTEIN\_MODIFICATION\_PROCESS | 3.76E-07 |
| GO\_SECRETORY\_VESICLE | 4.02E-07 |
| GO\_CARBOHYDRATE\_DERIVATIVE\_BIOSYNTHETIC\_PROCESS | 4.29E-07 |
| GO\_REGULATION\_OF\_CELLULAR\_COMPONENT\_SIZE | 4.83E-07 |
| GO\_LIPID\_BINDING | 5.23E-07 |
| GO\_MULTICELLULAR\_ORGANISM\_REPRODUCTION | 5.27E-07 |
| GO\_VOLTAGE\_GATED\_ION\_CHANNEL\_ACTIVITY | 5.35E-07 |
| GO\_NEGATIVE\_REGULATION\_OF\_CELL\_PROLIFERATION | 6.40E-07 |
| GO\_ACTIN\_BASED\_CELL\_PROJECTION | 7.03E-07 |
| KEGG\_CALCIUM\_SIGNALING\_PATHWAY | 7.23E-07 |
| GO\_EPITHELIAL\_CELL\_DIFFERENTIATION | 7.25E-07 |
| GO\_RESPONSE\_TO\_LIPID | 7.80E-07 |
| GO\_GROWTH | 8.00E-07 |
| GO\_POSITIVE\_REGULATION\_OF\_ION\_TRANSPORT | 8.01E-07 |
| GO\_DIVALENT\_INORGANIC\_CATION\_HOMEOSTASIS | 8.05E-07 |
| GO\_LEUKOCYTE\_ACTIVATION | 8.06E-07 |
| GO\_REGULATION\_OF\_CYTOKINE\_PRODUCTION | 8.45E-07 |
| GO\_NEUROTRANSMITTER\_TRANSPORT | 9.04E-07 |
| GO\_RESPONSE\_TO\_BIOTIC\_STIMULUS | 1.05E-06 |
| GO\_MONOVALENT\_INORGANIC\_CATION\_TRANSPORT | 1.05E-06 |
| GO\_PEPTIDASE\_ACTIVITY | 1.05E-06 |
| GO\_REGULATION\_OF\_PROTEOLYSIS | 1.06E-06 |
| GO\_HINDBRAIN\_DEVELOPMENT | 1.09E-06 |
| REACTOME\_HEMOSTASIS | 1.10E-06 |
| GO\_NEGATIVE\_REGULATION\_OF\_PHOSPHORUS\_METABOLIC\_PROCESS | 1.23E-06 |
| GO\_CELL\_RECOGNITION | 1.24E-06 |
| GO\_EXTRACELLULAR\_LIGAND\_GATED\_ION\_CHANNEL\_ACTIVITY | 1.25E-06 |
| GO\_REGULATION\_OF\_HOMEOSTATIC\_PROCESS | 1.27E-06 |
| GO\_HEART\_PROCESS | 1.44E-06 |
| GO\_CYTOPLASMIC\_VESICLE\_PART | 1.48E-06 |
| GO\_VASCULATURE\_DEVELOPMENT | 1.51E-06 |
| GO\_PRESYNAPTIC\_PROCESS\_INVOLVED\_IN\_SYNAPTIC\_TRANSMISSION | 1.89E-06 |
| REACTOME\_IMMUNE\_SYSTEM | 1.98E-06 |
| REACTOME\_TRANSMISSION\_ACROSS\_CHEMICAL\_SYNAPSES | 2.03E-06 |
| GO\_CELL\_CELL\_JUNCTION | 2.04E-06 |
| GO\_APICAL\_PLASMA\_MEMBRANE | 2.36E-06 |
| GO\_NEGATIVE\_REGULATION\_OF\_CELL\_DEVELOPMENT | 2.43E-06 |
| GO\_MULTI\_ORGANISM\_REPRODUCTIVE\_PROCESS | 2.49E-06 |
| GO\_REGULATION\_OF\_INTRACELLULAR\_TRANSPORT | 2.65E-06 |
| GO\_POSITIVE\_REGULATION\_OF\_CELL\_DEATH | 2.76E-06 |
| GO\_POSITIVE\_REGULATION\_OF\_IMMUNE\_RESPONSE | 2.77E-06 |
| GO\_MONOVALENT\_INORGANIC\_CATION\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY | 2.89E-06 |
| GO\_DEVELOPMENTAL\_PROCESS\_INVOLVED\_IN\_REPRODUCTION | 2.90E-06 |
| GO\_SYNAPSE\_ORGANIZATION | 3.36E-06 |
| GO\_REGULATION\_OF\_CALCIUM\_ION\_TRANSPORT | 3.50E-06 |
| GO\_EXTRACELLULAR\_STRUCTURE\_ORGANIZATION | 3.54E-06 |
| REACTOME\_GABA\_A\_RECEPTOR\_ACTIVATION | 3.65E-06 |
| GO\_DEVELOPMENTAL\_GROWTH | 3.86E-06 |
| GO\_REGULATION\_OF\_CELL\_ACTIVATION | 4.26E-06 |
| GO\_POSITIVE\_REGULATION\_OF\_LOCOMOTION | 4.41E-06 |
| GO\_RESPONSE\_TO\_OXYGEN\_LEVELS | 5.13E-06 |
| GO\_CELL\_PROJECTION\_MEMBRANE | 5.77E-06 |
| GO\_SULFUR\_COMPOUND\_BINDING | 5.93E-06 |
| GO\_REGULATION\_OF\_SEQUESTERING\_OF\_CALCIUM\_ION | 5.95E-06 |
| GO\_CARDIAC\_MUSCLE\_CELL\_CONTRACTION | 6.08E-06 |
| GO\_TUBE\_DEVELOPMENT | 6.25E-06 |
| REACTOME\_CELL\_CELL\_COMMUNICATION | 6.68E-06 |
| REACTOME\_AXON\_GUIDANCE | 6.73E-06 |
| GO\_VOCALIZATION\_BEHAVIOR | 6.74E-06 |
| GO\_TRANSMEMBRANE\_RECEPTOR\_PROTEIN\_PHOSPHATASE\_ACTIVITY | 6.89E-06 |
| GO\_RAS\_GUANYL\_NUCLEOTIDE\_EXCHANGE\_FACTOR\_ACTIVITY | 6.93E-06 |
| GO\_GABA\_RECEPTOR\_COMPLEX | 7.36E-06 |
| GO\_ACTIN\_FILAMENT\_BASED\_MOVEMENT | 7.84E-06 |
| GO\_POSITIVE\_REGULATION\_OF\_MAPK\_CASCADE | 7.88E-06 |
| GO\_POSITIVE\_REGULATION\_OF\_CELLULAR\_COMPONENT\_BIOGENESIS | 8.14E-06 |
| GO\_PROTEIN\_OLIGOMERIZATION | 8.24E-06 |
| GO\_REGULATION\_OF\_CYTOSKELETON\_ORGANIZATION | 8.70E-06 |
| GO\_ANION\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY | 8.71E-06 |
| GO\_EXTERNAL\_SIDE\_OF\_PLASMA\_MEMBRANE | 8.77E-06 |
| GO\_ION\_CHANNEL\_BINDING | 9.82E-06 |
| GO\_ACTIVATION\_OF\_IMMUNE\_RESPONSE | 9.92E-06 |
| GO\_REGULATION\_OF\_CYTOPLASMIC\_TRANSPORT | 1.06E-05 |
| GO\_REGULATION\_OF\_HEART\_RATE | 1.07E-05 |
| GO\_REGULATION\_OF\_DENDRITE\_DEVELOPMENT | 1.09E-05 |
| GO\_REGULATION\_OF\_HEART\_RATE\_BY\_CARDIAC\_CONDUCTION | 1.15E-05 |
| GO\_NEGATIVE\_REGULATION\_OF\_HYDROLASE\_ACTIVITY | 1.19E-05 |
| GO\_SYNAPSE\_ASSEMBLY | 1.19E-05 |
| GO\_CELL\_PROLIFERATION | 1.22E-05 |
| GO\_SEXUAL\_REPRODUCTION | 1.23E-05 |
| GO\_LYTIC\_VACUOLE | 1.27E-05 |
| GO\_SERINE\_HYDROLASE\_ACTIVITY | 1.28E-05 |
| GO\_NEGATIVE\_REGULATION\_OF\_CELL\_ADHESION | 1.30E-05 |
| GO\_SIGNAL\_RELEASE | 1.32E-05 |
| GO\_POSITIVE\_REGULATION\_OF\_CELL\_ADHESION | 1.32E-05 |
| GO\_ACTIN\_MEDIATED\_CELL\_CONTRACTION | 1.34E-05 |
| GO\_VOLTAGE\_GATED\_CATION\_CHANNEL\_ACTIVITY | 1.37E-05 |
| GO\_REGULATION\_OF\_SYNAPSE\_STRUCTURE\_OR\_ACTIVITY | 1.37E-05 |
| GO\_IMMUNE\_EFFECTOR\_PROCESS | 1.39E-05 |
| GO\_PHOSPHOLIPID\_BINDING | 1.39E-05 |
| GO\_DETECTION\_OF\_STIMULUS | 1.45E-05 |
| GO\_DENDRITE\_DEVELOPMENT | 1.53E-05 |
| GO\_STRUCTURAL\_MOLECULE\_ACTIVITY | 1.54E-05 |
| GO\_PEPTIDYL\_AMINO\_ACID\_MODIFICATION | 1.57E-05 |
| KEGG\_AXON\_GUIDANCE | 1.59E-05 |
| GO\_CELLULAR\_RESPONSE\_TO\_CYTOKINE\_STIMULUS | 1.63E-05 |
| GO\_CELL\_SUBSTRATE\_JUNCTION | 1.70E-05 |
| GO\_REGULATION\_OF\_CALCIUM\_ION\_IMPORT | 1.74E-05 |
| REACTOME\_LIGAND\_GATED\_ION\_CHANNEL\_TRANSPORT | 1.78E-05 |
| GO\_EAR\_DEVELOPMENT | 1.92E-05 |
| KEGG\_VASCULAR\_SMOOTH\_MUSCLE\_CONTRACTION | 2.00E-05 |
| GO\_RESPONSE\_TO\_GROWTH\_FACTOR | 2.01E-05 |
| GO\_EYE\_DEVELOPMENT | 2.05E-05 |
| GO\_POSITIVE\_REGULATION\_OF\_CELL\_MORPHOGENESIS\_INVOLVED\_IN\_DIFFERENTIATION | 2.13E-05 |
| GO\_CELL\_CYCLE\_PROCESS | 2.14E-05 |
| GO\_ADULT\_BEHAVIOR | 2.15E-05 |
| GO\_NITROGEN\_COMPOUND\_TRANSPORT | 2.19E-05 |
| GO\_NEURON\_CELL\_CELL\_ADHESION | 2.28E-05 |
| GO\_GABA\_RECEPTOR\_ACTIVITY | 2.33E-05 |
| GO\_GLYCOSAMINOGLYCAN\_BINDING | 2.38E-05 |
| GO\_GLAND\_DEVELOPMENT | 2.43E-05 |
| REACTOME\_NEUROTRANSMITTER\_RECEPTOR\_BINDING\_AND\_DOWNSTREAM\_TRANSMISSION\_IN\_THE\_POSTSYNAPTIC\_CELL | 2.48E-05 |
| GO\_REGULATION\_OF\_POTASSIUM\_ION\_TRANSMEMBRANE\_TRANSPORT | 2.59E-05 |
| GO\_CELL\_CELL\_SIGNALING\_INVOLVED\_IN\_CARDIAC\_CONDUCTION | 2.60E-05 |
| GO\_ACTION\_POTENTIAL | 2.72E-05 |
| GO\_NEGATIVE\_REGULATION\_OF\_LOCOMOTION | 2.78E-05 |
| GO\_EMBRYO\_DEVELOPMENT\_ENDING\_IN\_BIRTH\_OR\_EGG\_HATCHING | 2.94E-05 |
| GO\_REGULATION\_OF\_MULTI\_ORGANISM\_PROCESS | 2.96E-05 |
| GO\_REGULATION\_OF\_RELEASE\_OF\_SEQUESTERED\_CALCIUM\_ION\_INTO\_CYTOSOL\_BY\_SARCOPLASMIC\_RETICULUM | 2.96E-05 |
| GO\_REGULATION\_OF\_TRANSPORTER\_ACTIVITY | 2.98E-05 |
| GO\_REGULATION\_OF\_CARDIAC\_MUSCLE\_CONTRACTION | 3.05E-05 |
| GO\_REGULATION\_OF\_CELL\_GROWTH | 3.06E-05 |
| GO\_VACUOLAR\_PART | 3.14E-05 |
| GO\_RESPONSE\_TO\_INORGANIC\_SUBSTANCE | 3.17E-05 |
| GO\_NEUROMUSCULAR\_PROCESS | 3.18E-05 |
| GO\_CYTOSOLIC\_CALCIUM\_ION\_TRANSPORT | 3.23E-05 |
| GO\_REGULATION\_OF\_SYSTEMIC\_ARTERIAL\_BLOOD\_PRESSURE | 3.27E-05 |
| GO\_REGULATION\_OF\_RELEASE\_OF\_SEQUESTERED\_CALCIUM\_ION\_INTO\_CYTOSOL | 3.47E-05 |
| GO\_POSITIVE\_REGULATION\_OF\_ORGANELLE\_ORGANIZATION | 3.66E-05 |
| GO\_AGING | 3.68E-05 |
| GO\_GLYCOPROTEIN\_METABOLIC\_PROCESS | 3.78E-05 |
| GO\_COGNITION | 3.91E-05 |
| GO\_ORGANELLE\_LOCALIZATION | 4.10E-05 |
| GO\_EXOCYTOSIS | 4.25E-05 |
| GO\_REGULATION\_OF\_WNT\_SIGNALING\_PATHWAY | 4.61E-05 |
| GO\_LOCALIZATION\_WITHIN\_MEMBRANE | 4.85E-05 |
| GO\_REGULATION\_OF\_CATABOLIC\_PROCESS | 4.91E-05 |
| GO\_MEMBRANE\_MICRODOMAIN | 4.97E-05 |
| GO\_REGULATION\_OF\_CARDIAC\_CONDUCTION | 5.01E-05 |
| GO\_REGULATION\_OF\_NEUROTRANSMITTER\_LEVELS | 5.05E-05 |
| GO\_NEGATIVE\_REGULATION\_OF\_NERVOUS\_SYSTEM\_DEVELOPMENT | 5.21E-05 |
| GO\_SECRETORY\_GRANULE | 5.48E-05 |
| GO\_CELL\_COMMUNICATION\_INVOLVED\_IN\_CARDIAC\_CONDUCTION | 5.48E-05 |
| GO\_ANION\_TRANSMEMBRANE\_TRANSPORT | 5.54E-05 |
| GO\_CARDIAC\_MUSCLE\_CELL\_ACTION\_POTENTIAL | 5.92E-05 |
| GO\_PHOSPHORIC\_ESTER\_HYDROLASE\_ACTIVITY | 6.12E-05 |
| GO\_SCAFFOLD\_PROTEIN\_BINDING | 6.16E-05 |
| GO\_REGULATION\_OF\_PROTEIN\_SERINE\_THREONINE\_KINASE\_ACTIVITY | 6.46E-05 |
| GO\_PROTEIN\_SERINE\_THREONINE\_KINASE\_ACTIVITY | 6.47E-05 |
| GO\_VESICLE\_MEMBRANE | 6.47E-05 |
| GO\_CELL\_ADHESION\_MOLECULE\_BINDING | 6.49E-05 |
| GO\_CELLULAR\_RESPONSE\_TO\_ORGANIC\_CYCLIC\_COMPOUND | 6.90E-05 |
| GO\_POSITIVE\_REGULATION\_OF\_ENDOTHELIAL\_CELL\_PROLIFERATION | 6.91E-05 |
| GO\_CALMODULIN\_BINDING | 7.01E-05 |
| GO\_INNERVATION | 7.33E-05 |
| GO\_REGULATION\_OF\_STRIATED\_MUSCLE\_CONTRACTION | 8.05E-05 |
| GO\_NEURON\_MIGRATION | 8.20E-05 |
| GO\_NERVE\_DEVELOPMENT | 8.53E-05 |
| GO\_SECOND\_MESSENGER\_MEDIATED\_SIGNALING | 8.54E-05 |
| GO\_PLATELET\_ACTIVATION | 8.81E-05 |
| GO\_POTASSIUM\_ION\_TRANSPORT | 8.89E-05 |
| GO\_LYMPHOCYTE\_ACTIVATION | 9.22E-05 |
| GO\_POTASSIUM\_CHANNEL\_ACTIVITY | 9.23E-05 |
| GO\_VENTRICULAR\_CARDIAC\_MUSCLE\_CELL\_ACTION\_POTENTIAL | 9.29E-05 |
| GO\_ORGANIC\_ANION\_TRANSPORT | 9.40E-05 |
| GO\_CEREBELLAR\_CORTEX\_DEVELOPMENT | 9.48E-05 |
| GO\_HINDBRAIN\_MORPHOGENESIS | 9.62E-05 |