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Eye Morphogenesis (GO:0048592)
                                                                          Myoblast Differentiation (GO:0045445)
                                                   Regulation Of Skeletal Muscle Cell Differentiation (GO:2001014)
                                         Positive Regulation Of Epithelial To Mesenchymal Transition (GO:0010718)
                                                                  Extracellular Matrix Organization (GO:0030198)
                                                                       Collagen Fibril Organization (GO:0030199)
                                                    Positive Regulation Of Focal Adhesion Assembly (GO:0051894)
                                                             Regulation Of Hepatocyte Proliferation (GO:2000345)
                                          Leukocyte Chemotaxis Involved In Inflammatory Response (GO:0002232)
                                                        Negative Regulation Of Histone Acetylation (GO:0035067)
                                                          Negative Regulation Of Neuron Migration (GO:2001223)
                                                               Plasma Membrane Raft Organization (GO:0044857)
                                           Positive Regulation Of Cardiac Muscle Cell Differentiation (GO:2000727)
                                                                    Endodermal Cell Differentiation (GO:0035987)
                                                                              Microvillus Assembly (GO:0030033)
                                                               Extracellular Structure Organization (GO:0043062)
                                                     External Encapsulating Structure Organization (GO:0045229)
                                                    Notch Signaling Involved In Heart Development (GO:0061314)
                                                                   Rap Protein Signal Transduction (GO:0032486)
                                                    Negative Regulation Of Sprouting Angiogenesis (GO:1903671)
                                                         Regulation Of Circadian Sleep/Wake Cycle (GO:0042749)
                                                              Monosaccharide Biosynthetic Process (GO:0046364)
                                              Positive Regulation Of Oligodendrocyte Differentiation (GO:0048714)
                                                             Chondroitin Sulfate Metabolic Process (GO:0030204)
                                                                          Dendrite Self-Avoidance (GO:0070593)
                                                   Regulation Of Circadian Sleep/Wake Cycle, Sleep (GO:0045187)
                                                        Regulation Of Relaxation Of Cardiac Muscle (GO:1901897)
                                              Regulation Of Voltage-Gated Sodium Channel Activity (GO:1905150)
                                                  Dermatan Sulfate Proteoglycan Metabolic Process (GO:0050655)
                          Transmembrane Receptor Protein Tyrosine Phosphatase Signaling Pathway (GO:0007185)
                                                                 Morphogenesis Of An Endothelium (GO:0003159)
                                                       Negative Regulation Of Relaxation Of Muscle (GO:1901078)
                                                Positive Regulation Of Animal Organ Morphogenesis (GO:0110110)
                                                                   Endothelial Tube Morphogenesis (GO:0061154)
                                                                                 Myelin Assembly (GO:0032288)
                                                                  Cell-Substrate Junction Assembly (GO:0007044)
                                                                         Regulation Of Ossification (GO:0030278)
                                             Regulation Of Myosin-Light-Chain-Phosphatase Activity (GO:0035507)
                                                 Positive Regulation Of Chondrocyte Differentiation (GO:0032332)
                                                        Positive Regulation Of Kidney Development (GO:0090184)
                                               Regulation Of Mesenchymal Stem Cell Differentiation (GO:2000739)
                                                             Regulation Of Vascular Wound Healing (GO:0061043)
                                 Positive Regulation Of Cardiac Epithelial To Mesenchymal Transition (GO:0062043)
                                                              Regulation Of Muscle System Process (GO:0090257)
                                                                Ephrin Receptor Signaling Pathway (GO:0048013)
                          Regulation Of ER-associated Ubiquitin-Dependent Protein Catabolic Process (GO:1903069)
                                                      Positive Regulation Of Cartilage Development (GO:0061036)
                                                              Regulation Of Cartilage Development (GO:0061035)
                                                                  L-ascorbic Acid Metabolic Process (GO:0019852)
                                                         PERK-mediated Unfolded Protein Response (GO:0036499)
                                                     Regulation Of Peptidyl-Cysteine S-nitrosylation (GO:2000169)
                                                            Regulation Of Plasma Membrane Repair (GO:1905684) -
                              Negative Regulation Of Amyloid Precursor Protein Biosynthetic Process (GO:0042985)
                                              Negative Regulation Of Lens Fiber Cell Differentiation (GO:1902747) -
                                                                              Polyamine Transport (GO:0015846)
                                             Positive Regulation Of Cholesterol Biosynthetic Process (GO:0045542)
                                                  Positive Regulation Of Sterol Biosynthetic Process (GO:0106120)
                                                         Regulation Of Chondrocyte Differentiation (GO:0032330)
                                                                           Cytoplasmic Translation (GO:0002181)
                                                              Epithelial To Mesenchymal Transition (GO:0001837)
                                                                                 Gene Expression (GO:0010467)
                                                                                 mRNA Processing (GO:0006397)
                RNA Splicing, Via Transesterification Reactions With Bulged Adenosine As Nucleophile (GO:0000377)
                                                                  mRNA Splicing, Via Spliceosome (GO:0000398)
                              Cell Surface Receptor Signaling Pathway Involved In Cell-Cell Signaling (GO:1905114)
                                                                 CRD-mediated mRNA Stabilization (GO:0070934)
                                        Positive Regulation Of Non-Canonical Wnt Signaling Pathway (GO:2000052)
                                                                         Cell-Cell Signaling By Wnt (GO:0198738)
                                                     Regulation Of mRNA Splicing, Via Spliceosome (GO:0048024)
                                                                      Peptide Biosynthetic Process (GO:0043043)
                                                     Regulation Of Endodeoxyribonuclease Activity (GO:0032071)
                                                    Termination Of RNA Polymerase II Transcription (GO:0006369)
                                                                  Mesenchymal Cell Differentiation (GO:0048762)
                                                                                   RNA Processing (GO:0006396)
                                                                Smooth Muscle Cell Differentiation (GO:0051145)
Negative Regulation Of Nuclear-Transcribed mRNA Catabolic Process, Deadenylation-Dependent Decay (GO:1900152)
                                               Negative Regulation Of DNA-templated Transcription (GO:0045892)
            Positive Regulation Of Extrinsic Apoptotic Signaling Pathway Via Death Domain Receptors (GO:1902043)
                                                   Regulation Of Basement Membrane Organization (GO:0110011)
                                         Positive Regulation Of Integrin-Mediated Signaling Pathway (GO:2001046)
                                                 Regulation Of Integrin-Mediated Signaling Pathway (GO:2001044)
                                                                              Axonal Fasciculation (GO:0007413)
                                                       Negative Regulation Of Platelet Aggregation (GO:0090331)
                                                     Positive Regulation Of Glial Cell Differentiation (GO:0045687)
                        Negative Regulation Of Fibroblast Growth Factor Receptor Signaling Pathway (GO:0040037)
                                             Regulation Of Retrograde Transport, Endosome To Golgi (GO:1905279)
                                                                              Response To Heparin (GO:0071503) -
                                                                      Response To Manganese Ion (GO:0010042) -
                                                               Retinal Ganglion Cell Axon Guidance (GO:0031290) -
                                            Negative Regulation Of Excitatory Postsynaptic Potential (GO:0090394) -
                                                  Negative Regulation Of Trophoblast Cell Migration (GO:1901164) -
                                            Plus-End-Directed Organelle Transport Along Microtubule (GO:0072386)
                                                                                  Protein Sulfation (GO:0006477)
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0 1000 Combined Score

■ F4: DS_DPEP1+

■ F5: NGFR+

■ F4: DP HHIP+

■ F5: RAMP1+

F4: TNN+COCH+