Negative Regulation Of Monocyte Chemotaxis (GO:0090027) F1: Superficial Negative Regulation Of Cell Cycle Phase Transition (GO:1901988) F2: Universal Negative Regulation Of Mononuclear Cell Migration (GO:0071676) ■ F2/3: Perivascular Bleb Assembly (GO:0032060) Smooth Muscle Cell Migration (GO:0014909) ■ F3: FRC-like Nucleobase Catabolic Process (GO:0046113) Regulation Of Stem Cell Division (GO:2000035) Pyrimidine Nucleobase Catabolic Process (GO:0006208) Negative Regulation Of Leukocyte Chemotaxis (GO:0002689) Cyclooxygenase Pathway (GO:0019371) Pyrimidine-Containing Compound Catabolic Process (GO:0072529) Cellular Response To Glucocorticoid Stimulus (GO:0071385) Collagen Fibril Organization (GO:0030199) Extracellular Structure Organization (GO:0043062) Negative Regulation Of Wnt Signaling Pathway (GO:0030178) External Encapsulating Structure Organization (GO:0045229) Extracellular Matrix Organization (GO:0030198) Negative Regulation Of Canonical Wnt Signaling Pathway (GO:0090090) Sprouting Angiogenesis (GO:0002040) Regulation Of Focal Adhesion Disassembly (GO:0120182) Regulation Of Keratinocyte Apoptotic Process (GO:1902172) Regulation Of Long-Chain Fatty Acid Import Across Plasma Membrane (GO:0010746) Cellular Response To Granulocyte Macrophage Colony-Stimulating Factor Stimulus (GO:0097011) Response To Granulocyte Macrophage Colony-Stimulating Factor (GO:0097012) Mitochondrial Protein Catabolic Process (GO:0035694) Negative Regulation Of Erythrocyte Differentiation (GO:0045647) Negative Regulation Of Nervous System Process (GO:0031645) Photoreceptor Cell Differentiation (GO:0046530) Positive Regulation Of Wnt Signaling Pathway, Planar Cell Polarity Pathway (GO:2000096) Positive Regulation Of Extrinsic Apoptotic Signaling Pathway Via Death Domain Receptors (GO:1902043) Positive Regulation Of Focal Adhesion Disassembly (GO:0120183) Regulation Of Basement Membrane Organization (GO:0110011) Sequestering Of Extracellular Ligand From Receptor (GO:0035581) Response To Fibroblast Growth Factor (GO:0071774) Regulation Of Epithelial To Mesenchymal Transition (GO:0010717) Regulation Of Superoxide Metabolic Process (GO:0090322) Glycolipid Transport (GO:0046836) Elastic Fiber Assembly (GO:0048251) Maintenance Of Protein Location In Extracellular Region (GO:0071694) Contractile Actin Filament Bundle Assembly (GO:0030038) Stress Fiber Assembly (GO:0043149) Regulation Of Lipase Activity (GO:0060191) Mitochondrial Electron Transport, Cytochrome C To Oxygen (GO:0006123) Prostanoid Metabolic Process (GO:0006692) Regulation Of Peptidase Activity (GO:0052547) Cytoplasmic Translation (GO:0002181) Extracellular Matrix Assembly (GO:0085029) Regulation Of Neutrophil Activation (GO:1902563) Regulation Of Neutrophil Extravasation (GO:2000389) Regulation Of Plasma Membrane Repair (GO:1905684) Gamma-Aminobutyric Acid Metabolic Process (GO:0009448) Negative Regulation Of Cardiac Muscle Cell Apoptotic Process (GO:0010667) Negative Regulation Of Insulin-Like Growth Factor Receptor Signaling Pathway (GO:0043569) Positive Regulation Of Vesicle Fusion (GO:0031340) Protein Sulfation (GO:0006477) Cellular Response To Prostaglandin Stimulus (GO:0071379) Negative Regulation Of Establishment Of Protein Localization (GO:1904950) TORC1 Signaling (GO:0038202) Adherens Junction Maintenance (GO:0034334) Regulation Of Metallopeptidase Activity (GO:1905048) Angiotensin-Activated Signaling Pathway (GO:0038166) Definitive Hemopoiesis (GO:0060216) Ganglioside Catabolic Process (GO:0006689) Intracellular Sequestering Of Iron Ion (GO:0006880) Negative Regulation Of Membrane Protein Ectodomain Proteolysis (GO:0051045) Prostaglandin Metabolic Process (GO:0006693) Negative Regulation Of Actin Filament Polymerization (GO:0030837) Negative Regulation Of Interleukin-8 Production (GO:0032717) Positive Regulation Of Tau-Protein Kinase Activity (GO:1902949) Ribonucleoprotein Complex Biogenesis (GO:0022613) Peptide Biosynthetic Process (GO:0043043) Positive Regulation Of Cytoplasmic Translation (GO:2000767) Positive Regulation Of Protein Localization To Chromosome, Telomeric Region (GO:1904816) Gene Expression (GO:0010467) Ribosome Biogenesis (GO:0042254) Macromolecule Biosynthetic Process (GO:0009059) Regulation Of Transcription From RNA Polymerase II Promoter In Response To Stress (GO:0043618) Ribosomal Large Subunit Biogenesis (GO:0042273) Regulation Of Vascular Associated Smooth Muscle Cell Apoptotic Process (GO:1905459) Positive Regulation Of Establishment Of Protein Localization To Telomere (GO:1904851) Ribosomal Small Subunit Biogenesis (GO:0042274) Positive Regulation Of Signal Transduction By P53 Class Mediator (GO:1901798) Cellular Response To Arsenic-Containing Substance (GO:0071243) Positive Regulation Of Brown Fat Cell Differentiation (GO:0090336) Positive Regulation Of Transcription From RNA Polymerase II Promoter In Response To Stress (GO:0036003) CRD-mediated mRNA Stabilization (GO:0070934) Regulation Of Establishment Of Protein Localization To Telomere (GO:0070203) Regulation Of Nuclear-Transcribed mRNA Catabolic Process, Nonsense-Mediated Decay (GO:2000622) Regulation Of Protein Localization To Cajal Body (GO:1904869) Positive Regulation Of Protein Localization To Cajal Body (GO:1904871) Translation (GO:0006412) Regulation Of Transcription From RNA Polymerase II Promoter In Response To Hypoxia (GO:0061418) Protein Localization To Nucleolus (GO:1902570) Regulation Of Cellular Response To Heat (GO:1900034) Negative Regulation Of Nitrogen Compound Metabolic Process (GO:0051172) Negative Regulation Of Nuclear-Transcribed mRNA Catabolic Process, Deadenylation-Dependent Decay (GO:1900152) Negative Regulation Of Smooth Muscle Cell Proliferation (GO:0048662) Ribosome Assembly (GO:0042255) Positive Regulation Of Fat Cell Differentiation (GO:0045600) Response To Unfolded Protein (GO:0006986) Positive Regulation Of Cell Migration Involved In Sprouting Angiogenesis (GO:0090050) 'De Novo' Post-Translational Protein Folding (GO:0051084) Positive Regulation Of Monocyte Differentiation (GO:0045657) Positive Regulation Of Oligodendrocyte Differentiation (GO:0048714) Negative Regulation Of Cellular Extravasation (GO:0002692) Regulation Of Monocyte Differentiation (GO:0045655) Cellular Response To Interleukin-9 (GO:0071355) Vascular Associated Smooth Muscle Cell Development (GO:0097084) Interleukin-9-Mediated Signaling Pathway (GO:0038113) Regulation Of T-helper 1 Cell Differentiation (GO:0045625) Regulation Of T-helper Cell Differentiation (GO:0045622) Type II Interferon-Mediated Signaling Pathway (GO:0060333) Positive Regulation Of Macrophage Chemotaxis (GO:0010759) Regulation Of Erythrocyte Differentiation (GO:0045646) Response To Type II Interferon (GO:0034341) Cellular Response To Interleukin-15 (GO:0071350) Interleukin-15-Mediated Signaling Pathway (GO:0035723) Positive Regulation Of Macrophage Migration (GO:1905523) Response To Cytokine (GO:0034097) T-helper 17 Cell Lineage Commitment (GO:0072540) Negative Regulation Of Inflammatory Response To Wounding (GO:0106015) Negative Regulation Of Leukocyte Proliferation (GO:0070664) Positive Regulation Of CD4-positive, Alpha-Beta T Cell Differentiation (GO:0043372) Response To Interferon-Beta (GO:0035456) Cellular Response To Zinc Ion (GO:0071294) Regulation Of B Cell Activation (GO:0050864) Antigen Processing And Presentation Of Exogenous Peptide Antigen (GO:0002478) Positive Regulation Of Erythrocyte Differentiation (GO:0045648) Antigen Processing And Presentation Of Endogenous Peptide Antigen (GO:0002483) Cytokine-Mediated Signaling Pathway (GO:0019221) Cellular Response To Tumor Necrosis Factor (GO:0071356) T-helper 17 Cell Differentiation (GO:0072539) Endothelial Cell Chemotaxis (GO:0035767) Tyrosine Phosphorylation Of STAT Protein (GO:0007260) 0 500 750 1000 250 **Combined Score**