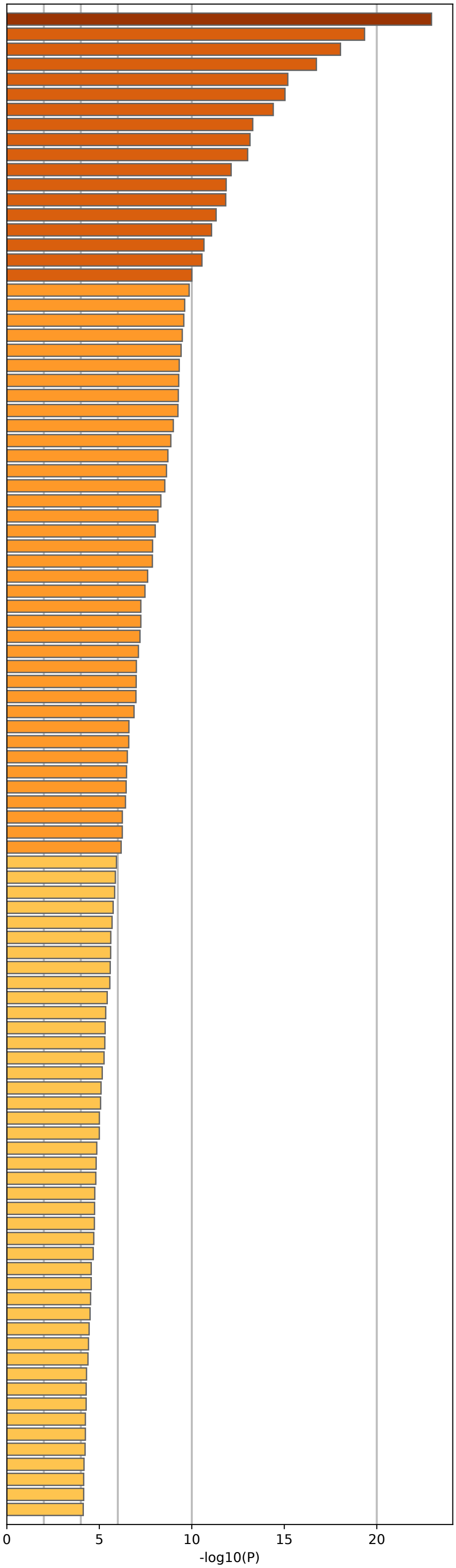

GO:0035239: tube morphogenesis
GO:0001501: skeletal system development
GO:0007507: heart development
M5884: NABA CORE MATRISOME
GO:0072001: renal system development
GO:0060485: mesenchyme development
GO:0009725: response to hormone
GO:0098609: cell-cell adhesion
GO:0061061: muscle structure development
R-HSA-1474244: Extracellular matrix organization
GO:0030198: extracellular matrix organization
GO:0007423: sensory organ development
GO:0090100: positive regulation of transmembrane receptor protein serine/threonine kinase signaling pathway
GO:0003013: circulatory system process
GO:0048732: gland development
hsa04350: TGF-beta signaling pathway
GO:0035107: appendage morphogenesis
GO:0050678: regulation of epithelial cell proliferation
GO:0035050: embryonic heart tube development
GO:0030855: epithelial cell differentiation
GO:0060348: bone development
M5885: NABA MATRISOME ASSOCIATED
GO:0071407: cellular response to organic cyclic compound
hsa05205: Proteoglycans in cancer
R-HSA-170834: Signaling by TGF-beta Receptor Complex
hsa04390: Hippo signaling pathway
GO:0048871: multicellular organismal-level homeostasis
GO:0045766: positive regulation of angiogenesis
GO:0042476: odontogenesis
GO:0030097: hemopoiesis
WP5087: Pleural mesothelioma
R-HSA-109582: Hemostasis
GO:0050878: regulation of body fluid levels
R-HSA-3000178: ECM proteoglycans
GO:0060541: respiratory system development
WP474: Endochondral ossification
GO:0030856: regulation of epithelial cell differentiation
GO:2000630: positive regulation of miRNA metabolic process
GO:0046942: carboxylic acid transport
GO:0001570: vasculogenesis
GO:1904888: cranial skeletal system development
GO:0030335: positive regulation of cell migration
GO:0043588: skin development
WP3967: miR 509 3p alteration of YAP1 ECM axis
GO:0032835: glomerulus development
GO:0071214: cellular response to abiotic stimulus
WP2857: Mesodermal commitment pathway
WP5090: Complement system in neuronal development and plasticity
WP236: Adipogenesis
M92: PID ANGIOPOIETIN RECEPTOR PATHWAY
GO:0040008: regulation of growth
GO:0098657: import into cell
GO:0010562: positive regulation of phosphorus metabolic process
GO:0007219: Notch signaling pathway
WP5144: NRP1 triggered signaling in pancreatic cancer
GO:0055123: digestive system development
GO:0045216: cell-cell junction organization
GO:0030111: regulation of Wnt signaling pathway
R-HSA-9006934: Signaling by Receptor Tyrosine Kinases
hsa04218: Cellular senescence
GO:0032570: response to progesterone
WP98: Prostaglandin synthesis and regulation
GO:0001656: metanephros development
GO:0043065: positive regulation of apoptotic process
GO:2000146: negative regulation of cell motility
hsa04810: Regulation of actin cytoskeleton
GO:0044057: regulation of system process
hsa04550: Signaling pathways regulating pluripotency of stem cells
GO:0051056: regulation of small GTPase mediated signal transduction
WP2034: Leptin signaling
GO:1903034: regulation of response to wounding
GO:0030278: regulation of ossification
GO:0030036: actin cytoskeleton organization
GO:1905686: positive regulation of plasma membrane repair
GO:0051145: smooth muscle cell differentiation
GO:0032350: regulation of hormone metabolic process
GO:0031099: regeneration
R-HSA-9012999: RHO GTPase cycle
WP5353: Macrophage stimulating protein MSP signaling
GO:0045936: negative regulation of phosphate metabolic process
GO:1990776: response to angiotensin
GO:0045087: innate immune response
R-HSA-2173782: Binding and Uptake of Ligands by Scavenger Receptors
GO:0048660: regulation of smooth muscle cell proliferation
GO:0007565: female pregnancy
GO:2000027: regulation of animal organ morphogenesis
WP3888: VEGFA VEGFR2 signaling
GO:0045598: regulation of fat cell differentiation
GO:0007548: sex differentiation
hsa04520: Adherens junction
GO:0045785: positive regulation of cell adhesion
WP5115: Network map of SARS CoV 2 signaling
GO:0072091: regulation of stem cell proliferation
GO:0051148: negative regulation of muscle cell differentiation
GO:0072089: stem cell proliferation
GO:0009299: mRNA transcription
R-HSA-9006931: Signaling by Nuclear Receptors
GO:0048368: lateral mesoderm development
GO:0090136: epithelial cell-cell adhesion
GO:0043255: regulation of carbohydrate biosynthetic process