A
Elbow plot


B


R-HSA-8953854: Metabolism of RNA
GO:0022613: ribonucleoprotein complex biogenesis
R-HSA-1640170: Cell Cycle
GO:0050684: regulation of mRNA processing
R-HSA-73856: RNA Polymerase II Transcription Termination
GO:0006259: DNA metabolic process
R-HSA-9716542: Signaling by Rho GTPases, Miro GTPases and RHOBTB3
R-HSA-1280215: Cytokine Signaling in Immune system
GO:0042254: ribosome biogenesis
GO:0006325: chromatin organization
R-HSA-72165: mRNA Splicing - Minor Pathway R-HSA-73894: DNA Repair
CORUM:3055: Na
CORUM:3055: Nop56p-associated pre-rRNA complex
R-HSA-3108232: SUMO E3 ligases SUMOylate target proteins
R-HSA-2262752: Cellular responses to stress
WP23: B cell receptor signaling pathway
R-HSA-168255: Influenza Infection
GO:0051052: regulation of DNA metabolic process GO:0006260: DNA replication
GO:0006401: RNA catabolic process



WP383: Striated muscle contraction pathway GO:0006941: striated muscle contraction M5884: NABA CORE MATRISOME R-HSA-114608: Platelet degranulation WP2118: Arrhythmogenic right ventricular cardiomyopathy hsa04610: Complement and coagulation cascades
GO:0030336: negative regulation of cell migration
GO:0030198: extracellular matrix organizatio
R-HSA-445355: Smooth Muscle Contraction
GO:0060538: skeletal muscle organ development
GO:0043268: positive regulation of potassium ion transport
GO:1902905: positive regulation of supramolecular fiber organization
WP2572: Primary focal segmental glomerulosclerosis (FSGS) WP5055: Burn wound healing
GO:0045765: regulation of angiogenesis
R-HSA-8978868: Fatty acid metabolism GO:0008217: regulation of blood pressure M3008: NABA ECM GLYCOPROTEINS


GO:0008544: epidermis development GO:0014706: striated muscle tissue development GO:0030335: positive regulation of cell migration

