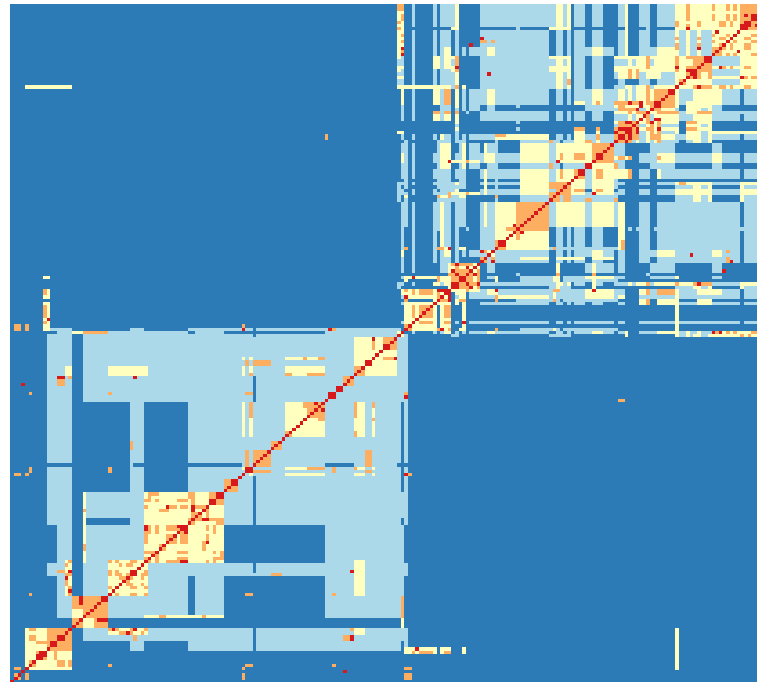
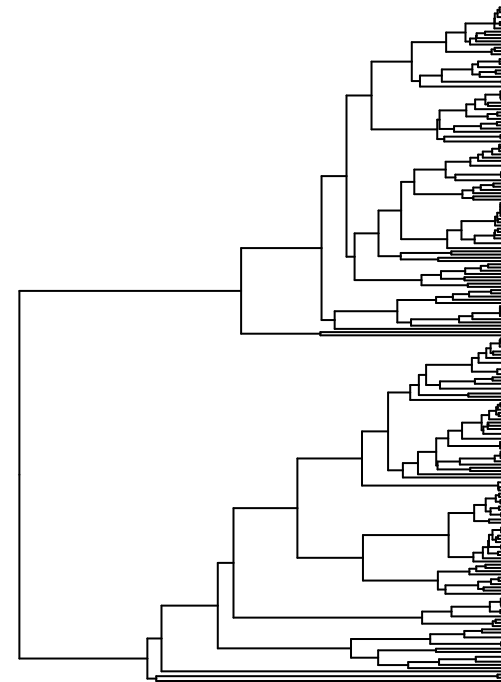
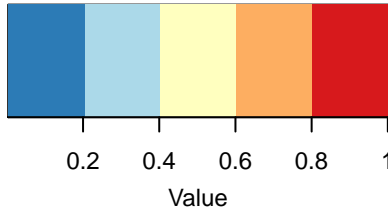


Color Key



GO:0042980: negative regulation of amyloid precursor protein biosynthetic process
GO:0045892: negative regulation of transcription, DNA-templated
GO:0042304: regulation of fatty acid biosynthetic process
GO:0004440: regulation of translational termination
GO:2001280: positive regulation of unsaturated fatty acid biosynthetic process
GO:0070495: negative regulation of thrombin-activated receptor signaling pathway
GO:1800408: negative regulation of cellular response to oxidative stress
GO:0048662: negative regulation of smooth muscle cell proliferation
GO:0070886: positive regulation of calcineurin-NFAT signaling cascade
GO:0045880: positive regulation of smoothened signaling pathway
GO:0050729: positive regulation of inflammatory response
GO:0010603: regulation of tumor necrosis factor-mediated signaling pathway
GO:0045088: regulation of innate immune response
GO:0035666: TRIF-dependent toll-like receptor signaling pathway
GO:1904179: positive regulation of adipose tissue development
GO:0045579: positive regulation of B cell differentiation
GO:0032725: positive regulation of interferon-gamma production
GO:0032823: regulation of natural killer cell differentiation
GO:0030502: negative regulation of bone mineralization
GO:0018098: negative regulation of relaxation of cardiac muscle
GO:0081347: planar cell polarity pathway involved in outflow tract morphogenesis
GO:0081354: planar cell polarity pathway involved in pericardium morphogenesis
GO:1904955: planar cell polarity pathway involved in midbrain dopaminergic neuron differentiation
GO:0007223: Wnt signaling pathway, calcium modulating pathway
GO:0018634: cAMP-mediated signaling
GO:0070245: positive regulation of thymocyte apoptotic process
GO:0032988: positive regulation of intracellular transport
GO:0032880: regulation of protein localization
GO:0006262: negative regulation of SMAD protein signal transduction
GO:1904457: positive regulation of neuronal action potential
GO:1900452: regulation of long-term synaptic depression
GO:0080648: membrane depolarization during bundle of His cell action potential
GO:1904199: positive regulation of regulation of vascular smooth muscle cell membrane depolarization
GO:0036494: positive regulation of translation initiation in response to endoplasmic reticulum stress
GO:0044700: oligodendrocyte differentiation
GO:0072560: type B pancreatic cell maturation
GO:0081028: establishment of endothelial barrier
GO:0001886: endothelial cell morphogenesis
GO:0021785: branchiomotor neuron axon guidance
GO:0018189: axon midline choice point recognition
GO:0042113: B cell activation
GO:0006029: convergent extension involved in organogenesis
GO:0035116: embryonic hindlimb morphogenesis
GO:0097094: craniofacial suture morphogenesis
GO:0060447: bud outgrowth involved in lung branching
GO:0007370: ventral furrow formation
GO:0000157: urinary bladder development
GO:0060740: prostate gland epithelium morphogenesis
GO:0007638: mechanosensory behavior
GO:1904954: sebaceous gland cell proliferation
GO:0046338: phosphatidylethanolamine catabolic process
GO:0038513: oxidative RNA demethylation
GO:0042775: mitochondrial ATP synthesis coupled electron transport
GO:0006351: transcription, DNA-templated
GO:0006025: mRNA cleavage involved in gene silencing by siRNA
GO:0031860: telomeric 3' overhang formation
GO:0042417: dopamine metabolic process
GO:0070830: bicellular tight junction assembly
GO:0014686: skeletal myofibril assembly
GO:0006698: postsynaptic assembly
GO:0030717: oocyte karyosome formation
GO:0071799: cellular response to prostaglandin D stimulus
GO:0070417: cellular response to cold
GO:0071774: response to fibroblast growth factor
GO:0035456: cargo loading into vesicle
GO:0015810: aspartate transmembrane transport
GO:0080609: membrane repolarization
GO:0051660: establishment of centrosome localization
GO:0086517: chemoattraction of serotonergic neuron axon
GO:0033627: cell adhesion mediated by integrin