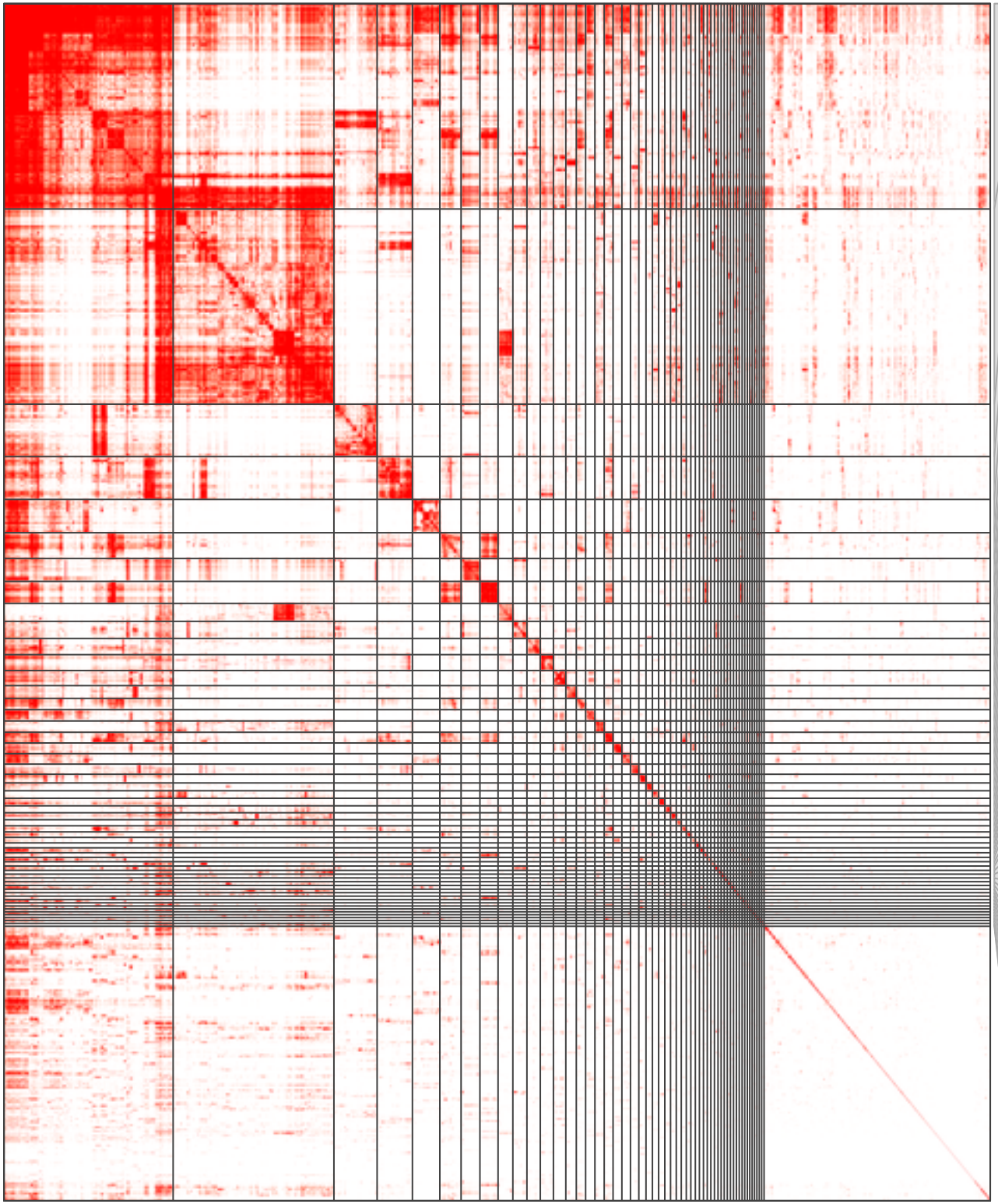
PND15_Adult−intergenic_bg_all_GO_BP



metabolic activity biosynthetic response
positive muscle
differentiation morphogenesis cardiac negative involved
negative membrane nucleus interleukin acid import
pathway alphabeta activation differentiation
negative amino positive catabolic
growth stimulus positive receptor chemical
process biosynthetic negative growth chemokine beta
pathway acid regulation receptor
negative
regulation mesenchymal differentiation kidney morphogenesis
secretion process hormone smooth system
involved signaling pathway negative death positive
leukocyte stem involved epithelial precursor
nuclear assembly centrosome
positive generation concentration calcium resorption bone
negative positive growth smad transduction signal
negative deacetylation remodeling acetylation peptidylarginine chromatin
involved transcription neuron spinal cord
response cell endothelial defense stimulus cellular mast
biogenesis endonucleolytic subunit ribonucleoprotein
catabolic sulfate positive glycoprotein heparan
leukocyte cellcell cellmatrix aggregation positive endothelial
transcription regulatory region receptor
feeding
secreting hypothalamus forebrain regulation neuron axonal
replication positive genome morphology
formation
pathway anteriorposterior regulation dorsalventral wnt
elongation positive morphogenesis muscle cell axis
cameratype optic formation morphogenesis otolith
eosinophil negative axon smooth muscle
polymerase hypoxia hyperoxia stimulus
inflammasome nucleation microtubule proteasome
cranial vestibulocochlear trigeminal
biogenic catabolic spermine spermidine
phorbol myristate acetate
monooxygenase ligase negative
arterial positive reninangiotensin involved negative circulatory
node system purkinje bundle
negative positive cardioblast mesenchymal neural crest
membrane maintenance myelin golgi regulation
positive
regulation catecholamine positive melanin
epithelial
pathway neural involved formation
signaling pathway folliclestimulating gonadotropinreleasing
myosinlightchainphosphatase phosphatase peptidyltyrosine peptidylserine
positive adenylate
regulation positive vascular development negative myoblast
compound sulfide hydrogen sulfation
gland
plasma highdensity remodeling positive
morphogenesis artery regulation vascular coronary development aorta

Similarity
0.3
0.2
0.1
0