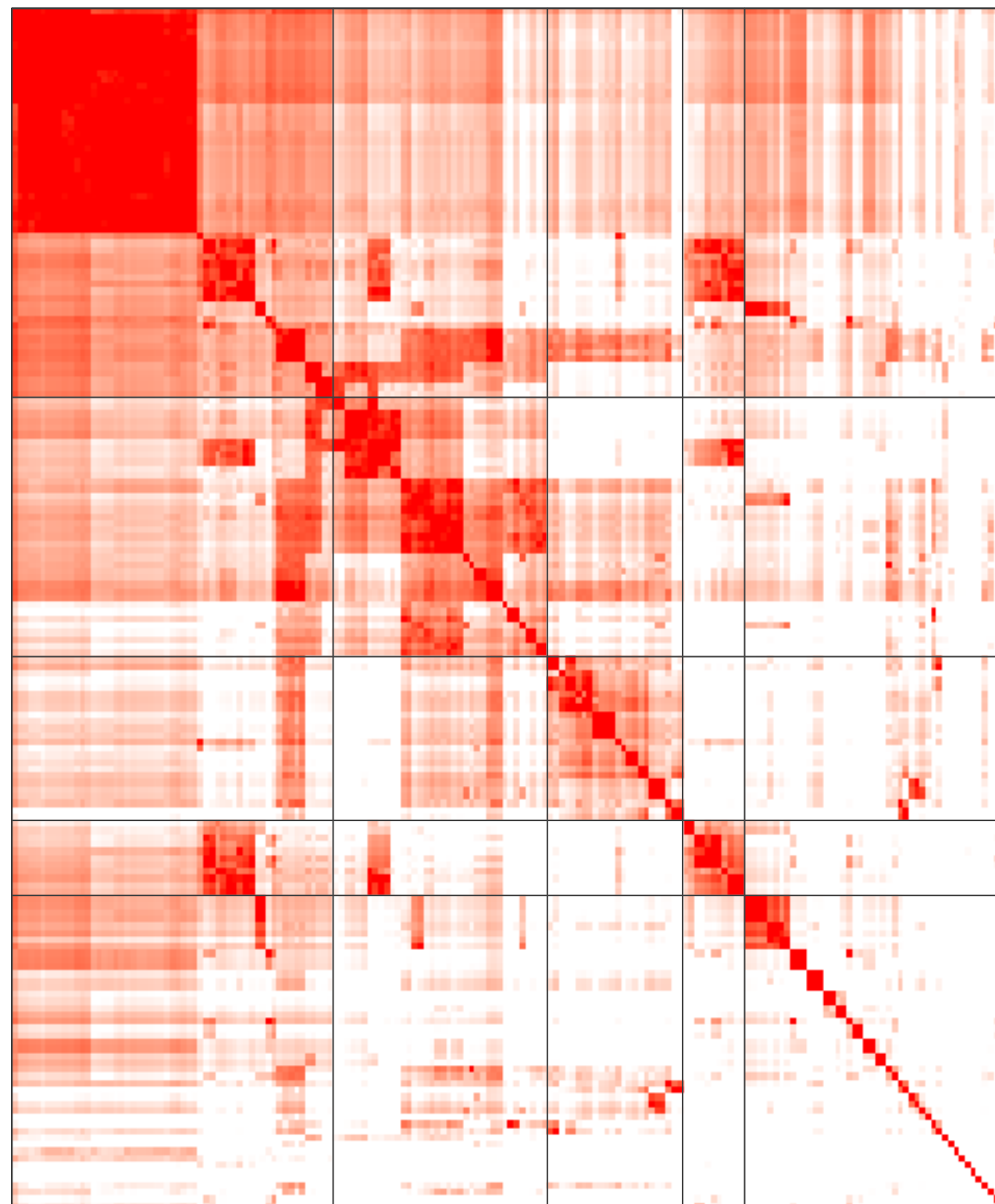


hIP_split_rna_cutoffproximal.active.accUp.rnaDown-proximal_2500bp-true_GO_BP



metabolic cell compound rna transcription
regulation process biosynthetic cellular positive

leukocyte negative immunity mast degranulation
regulation differentiation positive myeloid cell

spinal cord commitment maturation fate
development neuron cell forebrain differentiation

pathway engulfment
regulation positive phagocytosis uptake secretory regulated

Similarity
0.8
0.6
0.4
0.2
0