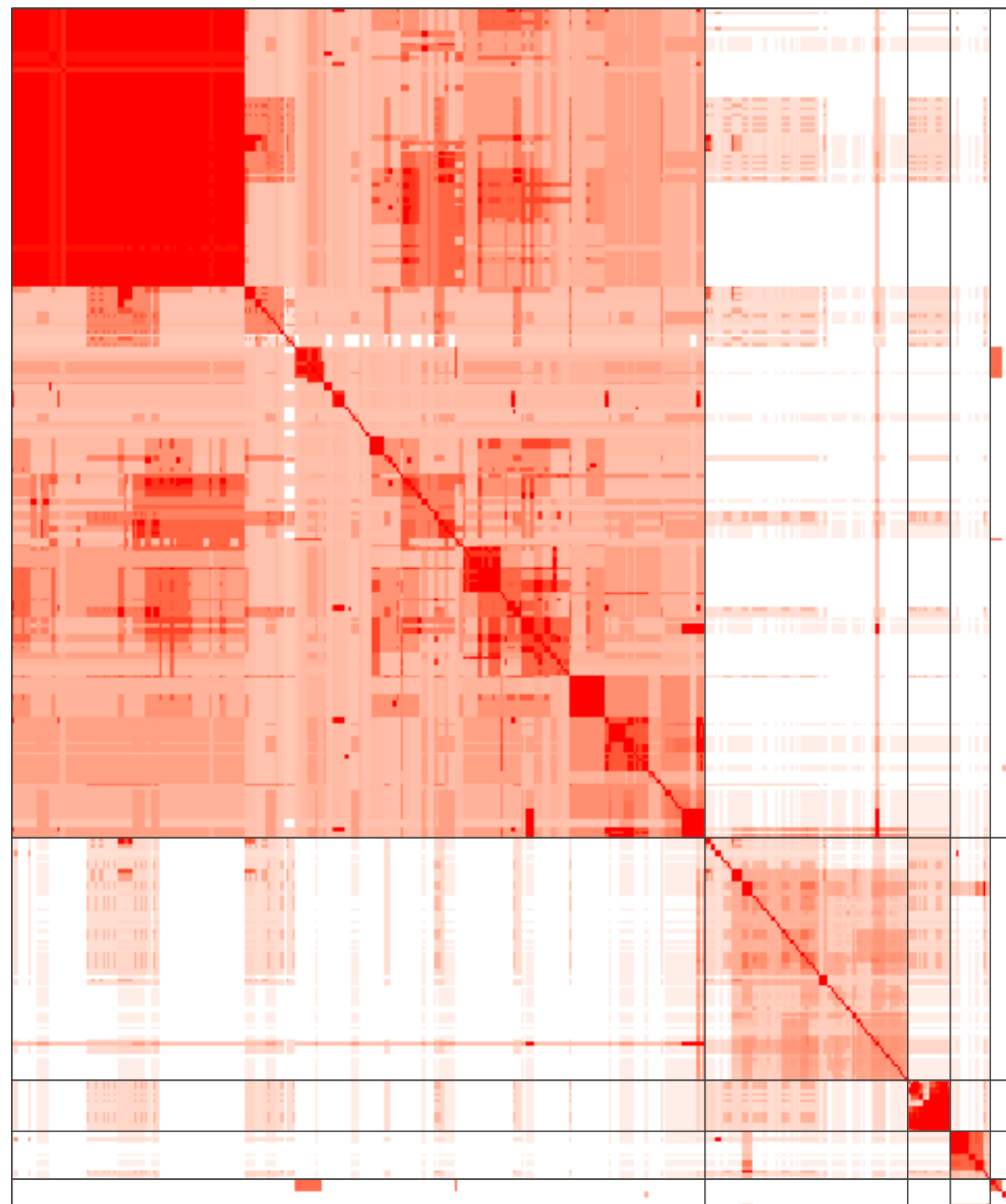


with_ChIP_split_proximal.inactive.accUp.H3K27me3-proximal_2500bp_GO_BP



differentiation cell involved formation tubule process metanephric
development morphogenesis regulation

Similarity
0.5
0.4
0.3
0.2
0.1
0

positive compound cellular rna transcription macromolecule
process regulation metabolic biosynthetic

cohesion division segregation chromosome maintenance
regulation sister positive chromatid mitotic

monosaccharide glucose peroxide hydrogen
response cellular stimulus retinoic acid hexose