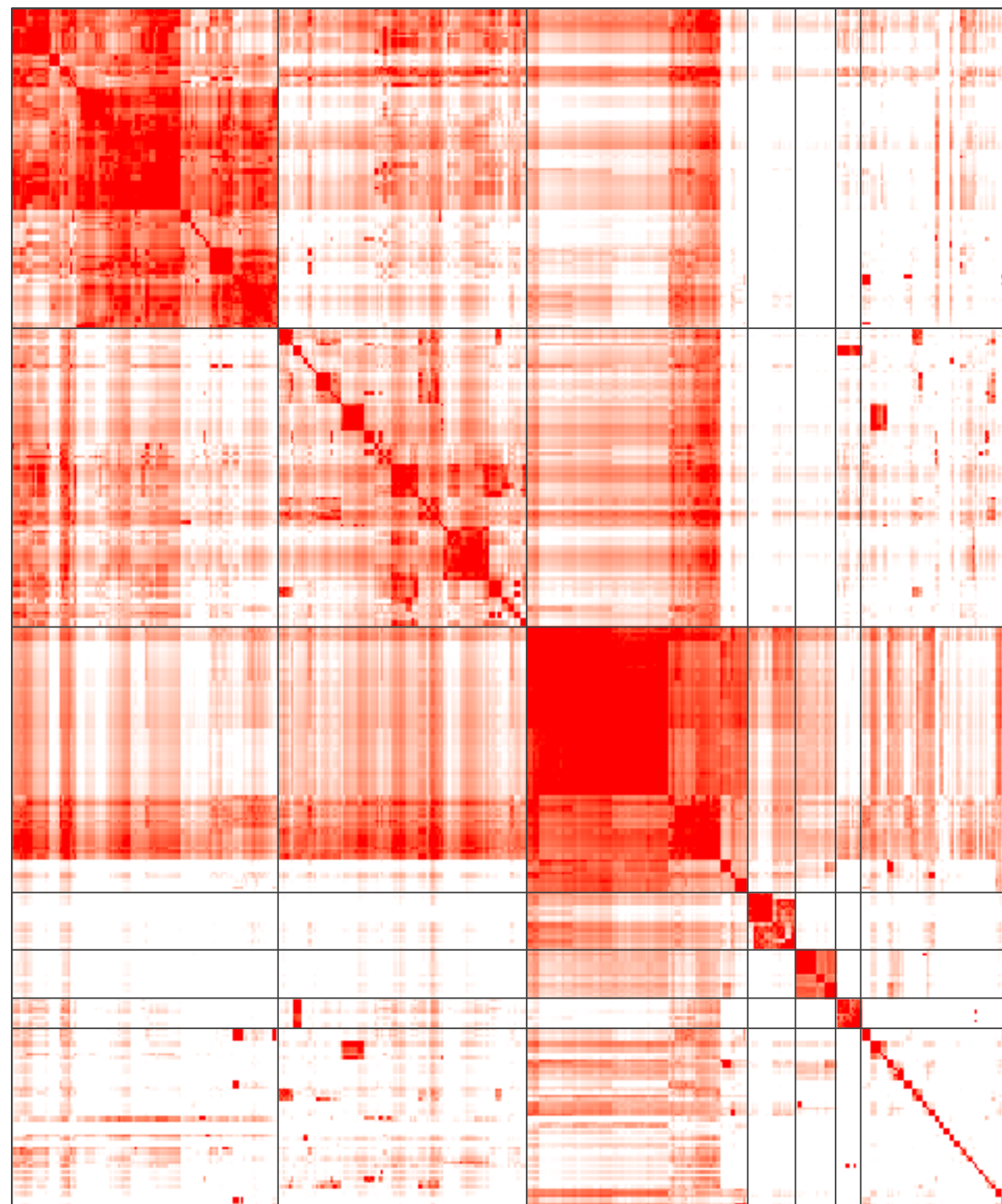


with_ChIP_proximal.inactive.accUp.H3K27me3_GO_BP



metanephric involved tubule kidney epithelial cell nephron
development regulation morphogenesis

differentiation embryonic cell neuron eye specification organ fate
development morphogenesis

positive cellular compound rna macromolecule transcription
process regulation metabolic biosynthetic

cohesion division segregation chromosome maintenance
regulation positive sister chromatid mitotic

glucose peroxide hydrogen
response cellular acid stimulus retinoic hexose monosaccharide

morphogenesis structural organization
nerve optic development vestibulocochlear formation cranial

Similarity
0.8
0.6
0.4
0.2
0