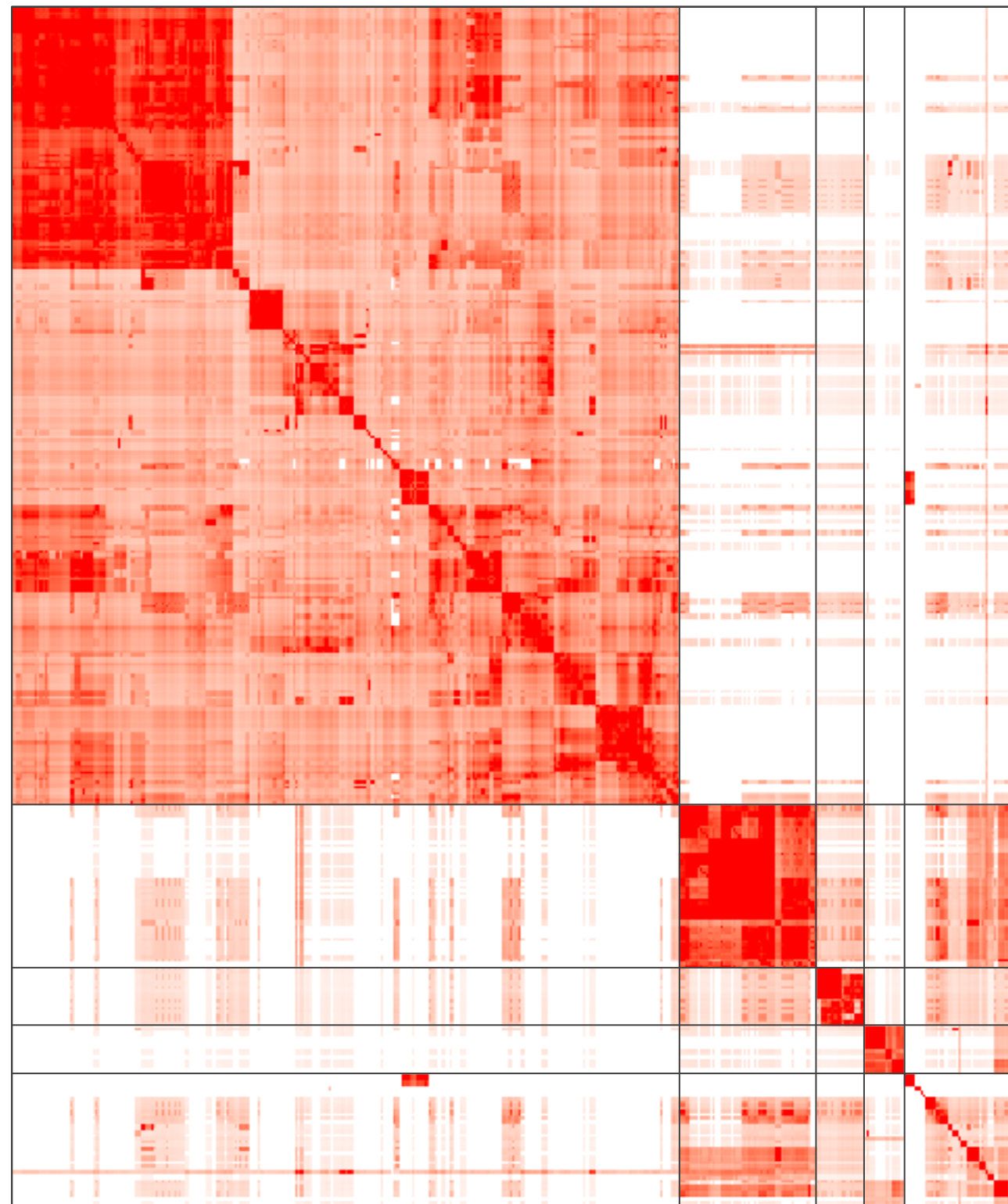


with\_ChIP\_proximal.inactive.accUp.H3K27me3\_GO\_BP



differentiation cell involved process tubule metanephric formation  
development morphogenesis regulation

compound positive cellular macromolecule transcription rna  
process regulation biosynthetic metabolic

cohesion division segregation chromosome maintenance  
regulation positive sister chromatid mitotic

glucose peroxide hydrogen  
response cellular acid stimulus retinoic hexose monosaccharide

Similarity  
0.8  
0.6  
0.4  
0.2  
0