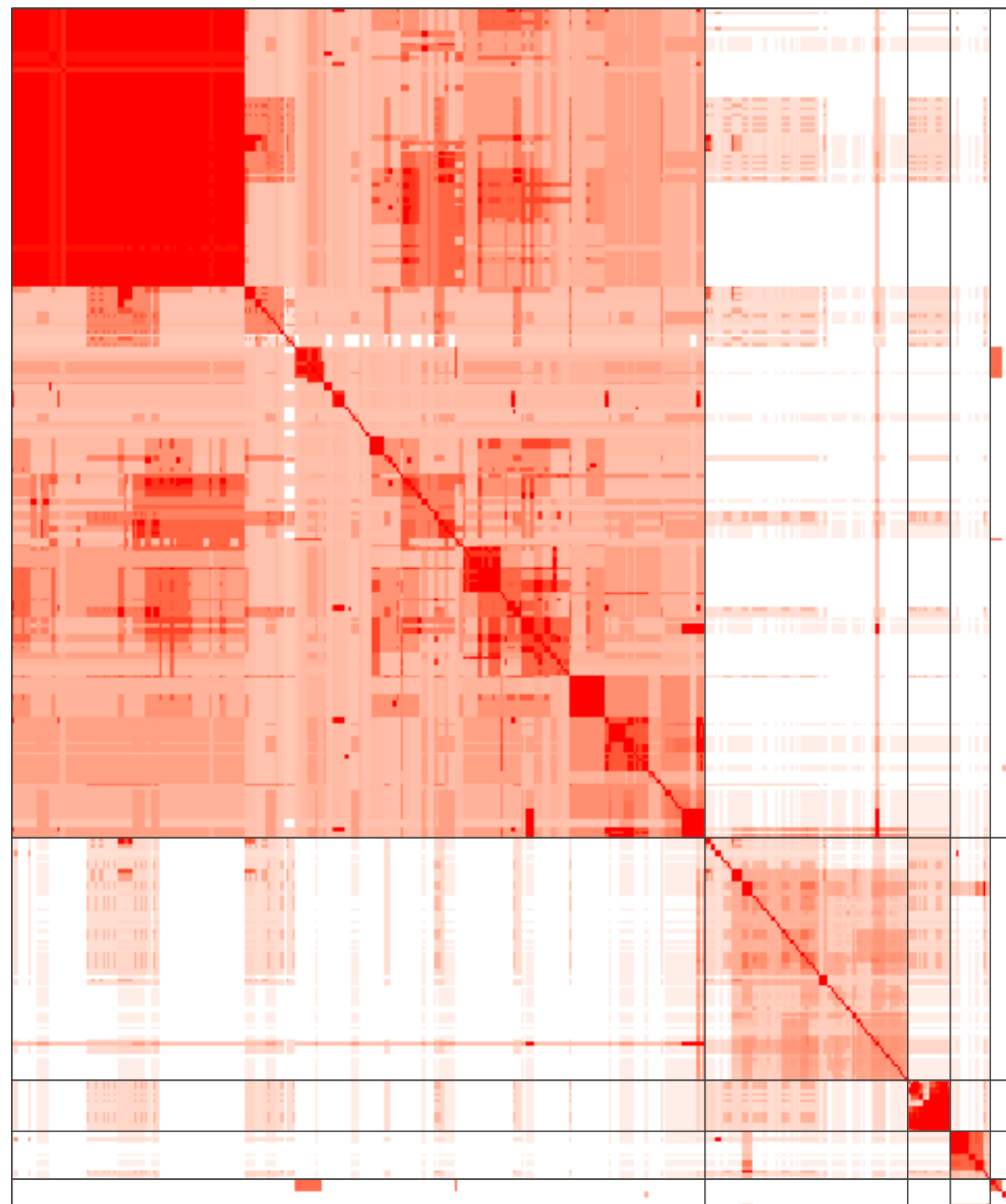


olig\_rna\_cutoff\_proximal.inactive.accUp.H3K27me3-proximal\_2500bp-not\_detected\_GO\_BP



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