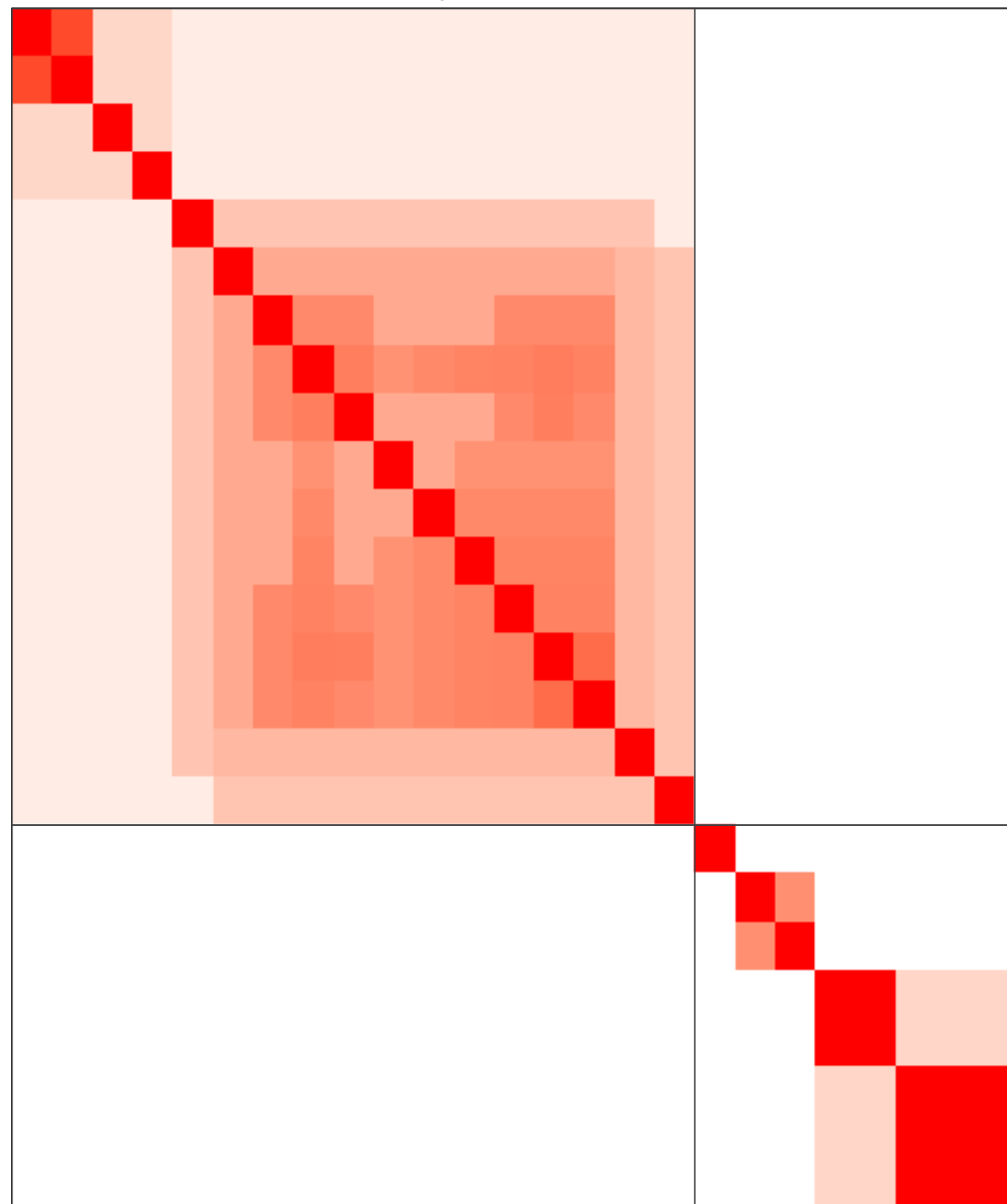


detected-with_ChIP_rna_cutoff_bg_all_proximal.inactive.accUp.H3K27me3_GO_MF



regulatory region promoter doublestranded
binding dna sequencespecific transcription rna polymerase

Similarity
0.6
0.4
0.2
0