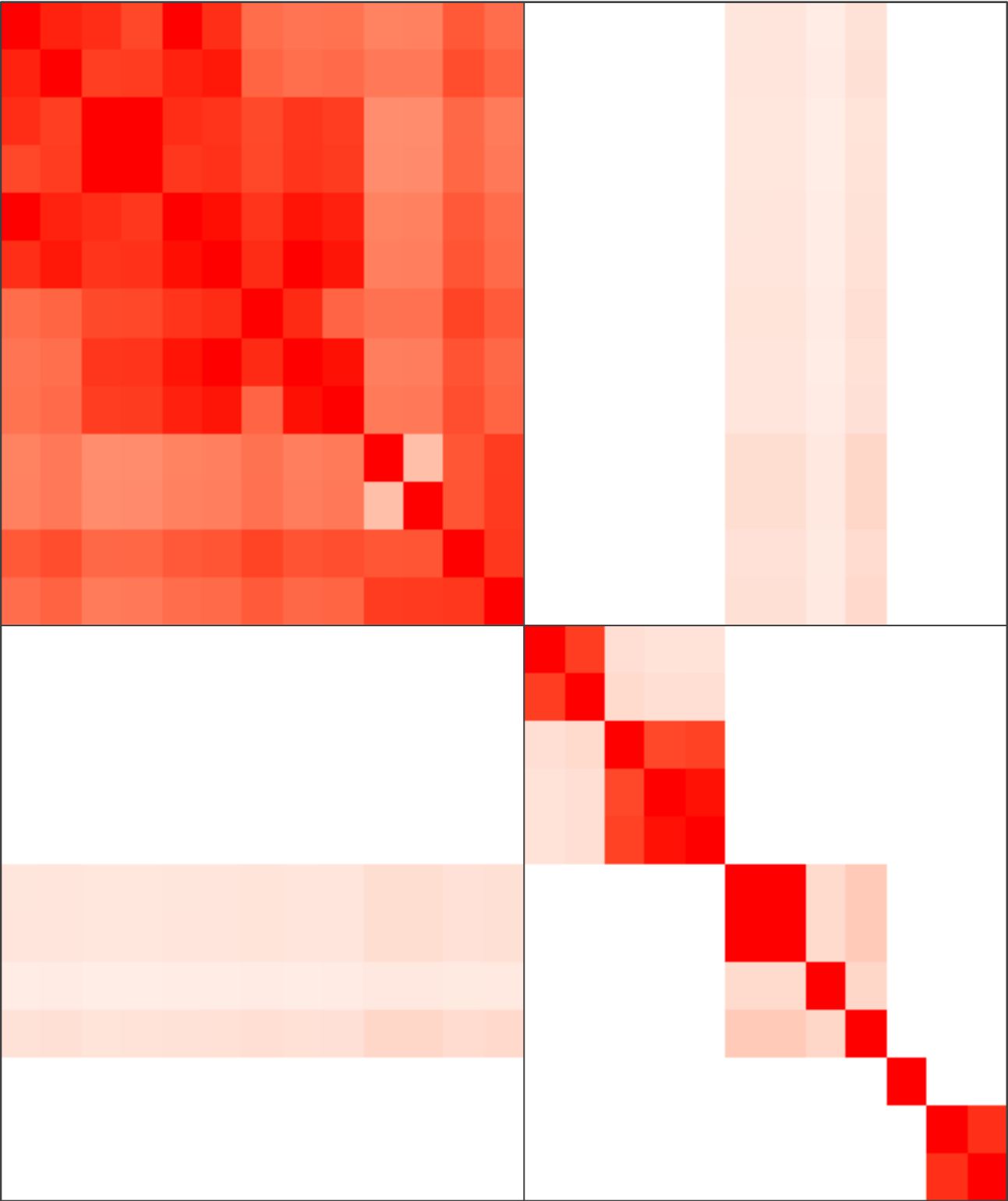


with_ChIP_bg_all_proximal.inactive.accUp.H3K27me3_GO_MF



polymerase promoter doublestranded transcription
binding dna sequencespecific regulatory region rna

Similarity
1
0.5
0