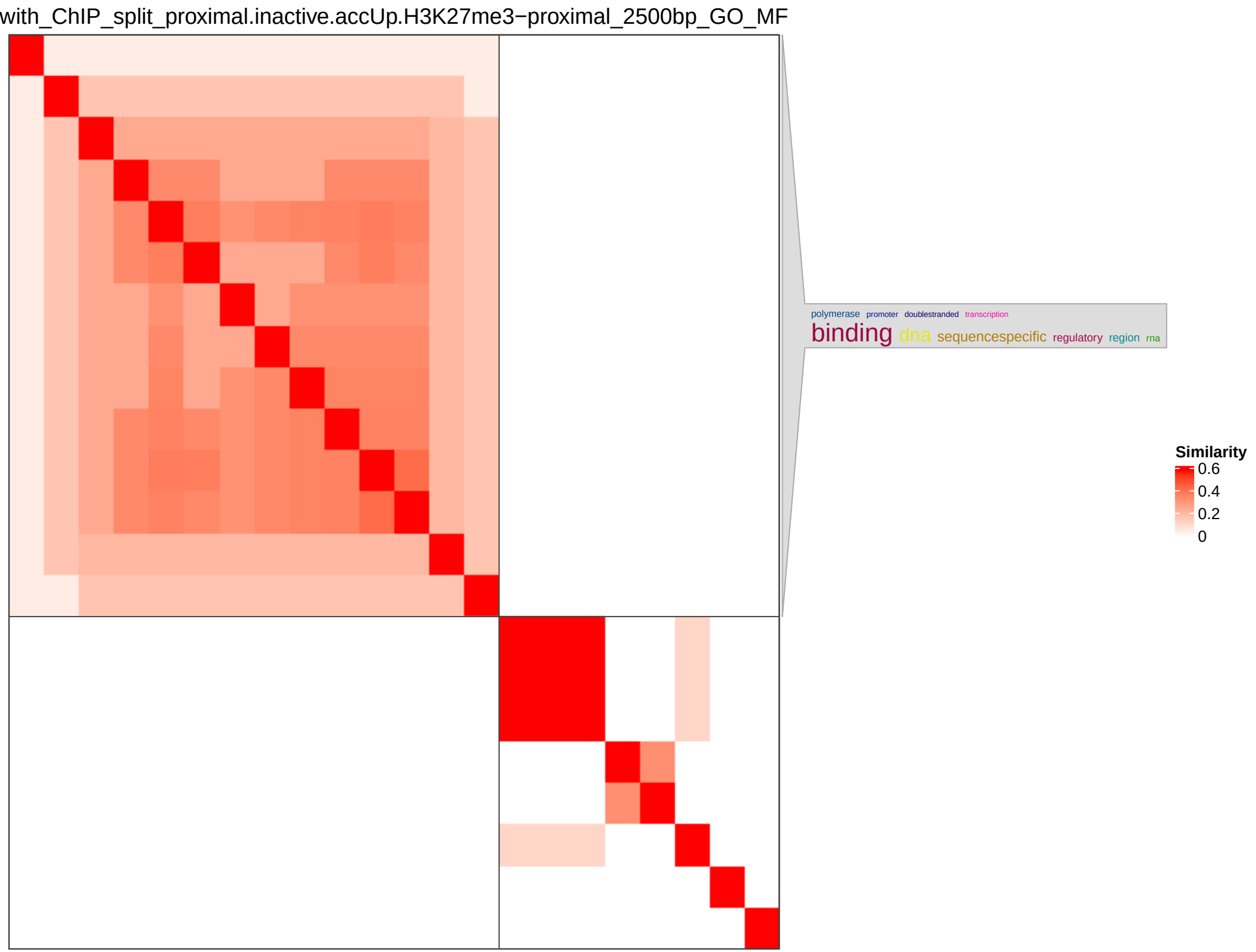
with_ChIP_split_proximal.inactive.accUp.H3K27me3−proximal_2500bp_GO_MF
polymerase promoter doublestranded transcription
binding dna sequencespecific regulatory region rna
Similarity
0.6
0.4
0.2
0