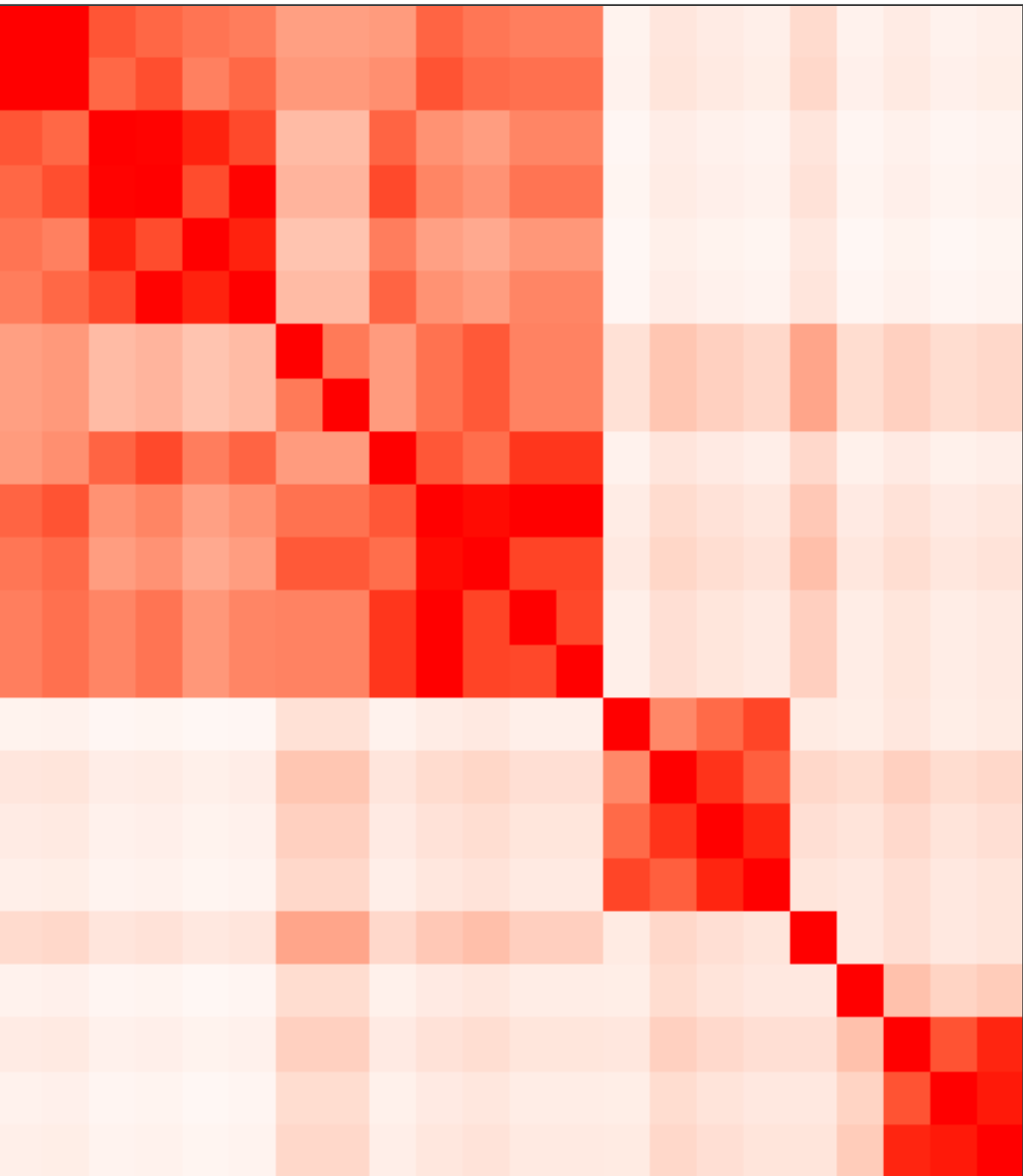


with_ChIP_split_proximal.inactive.accUp.H3K27me3-proximal_2500bp_GO_MF



Similarity

1
0.8
0.6
0.4
0.2
0

polymerase transcription promoter regulatory region
binding dna activity sequencespecific rna