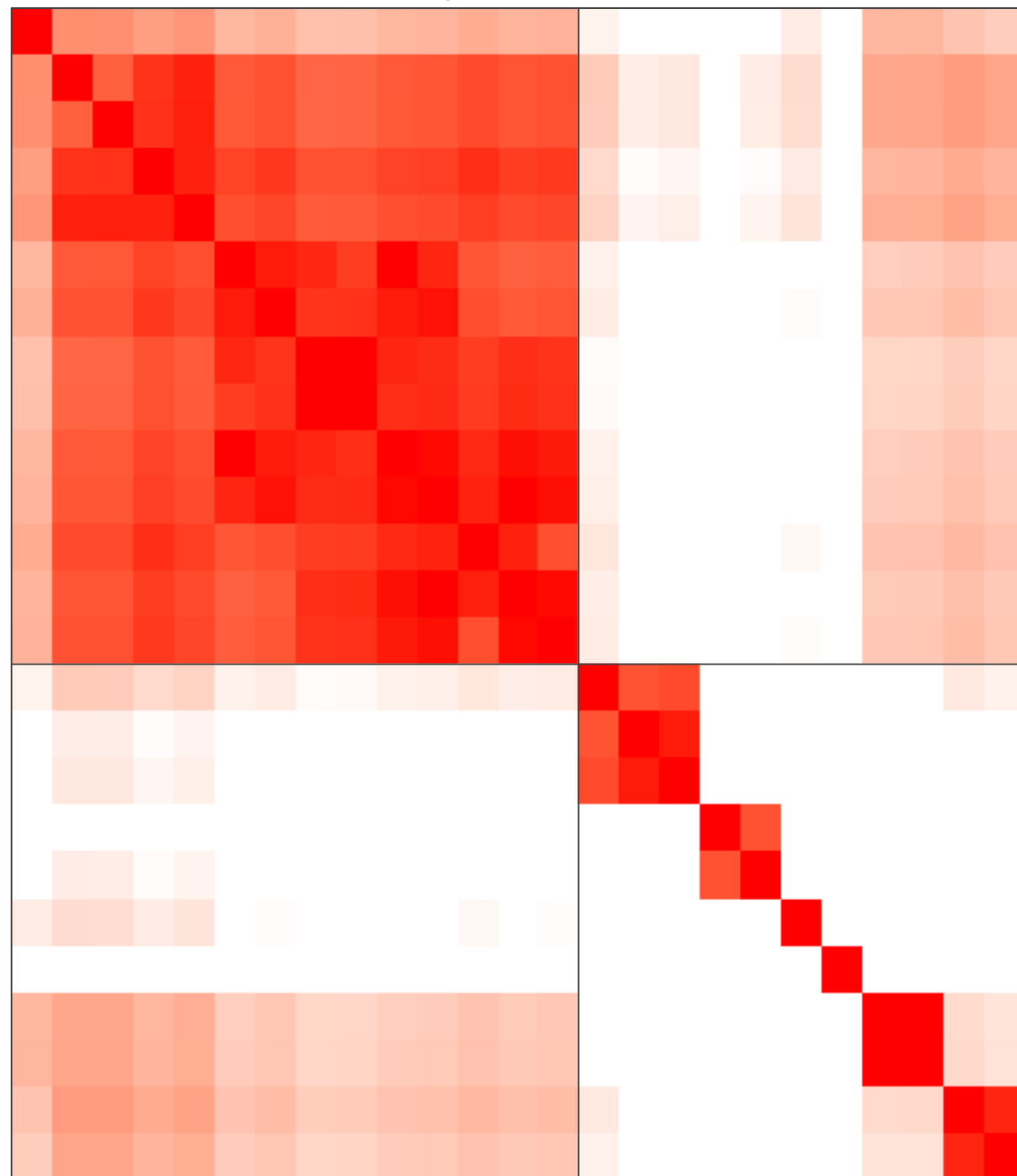


\_detected-with\_ChIP\_rna\_cutoff\_bg\_all\_proximal.inactive.accUp.H3K27me3\_GO\_MF



transcription rna polymerase promoter doublestranded  
binding dna sequencespecific regulatory region

**Similarity**  
1  
0.5  
0