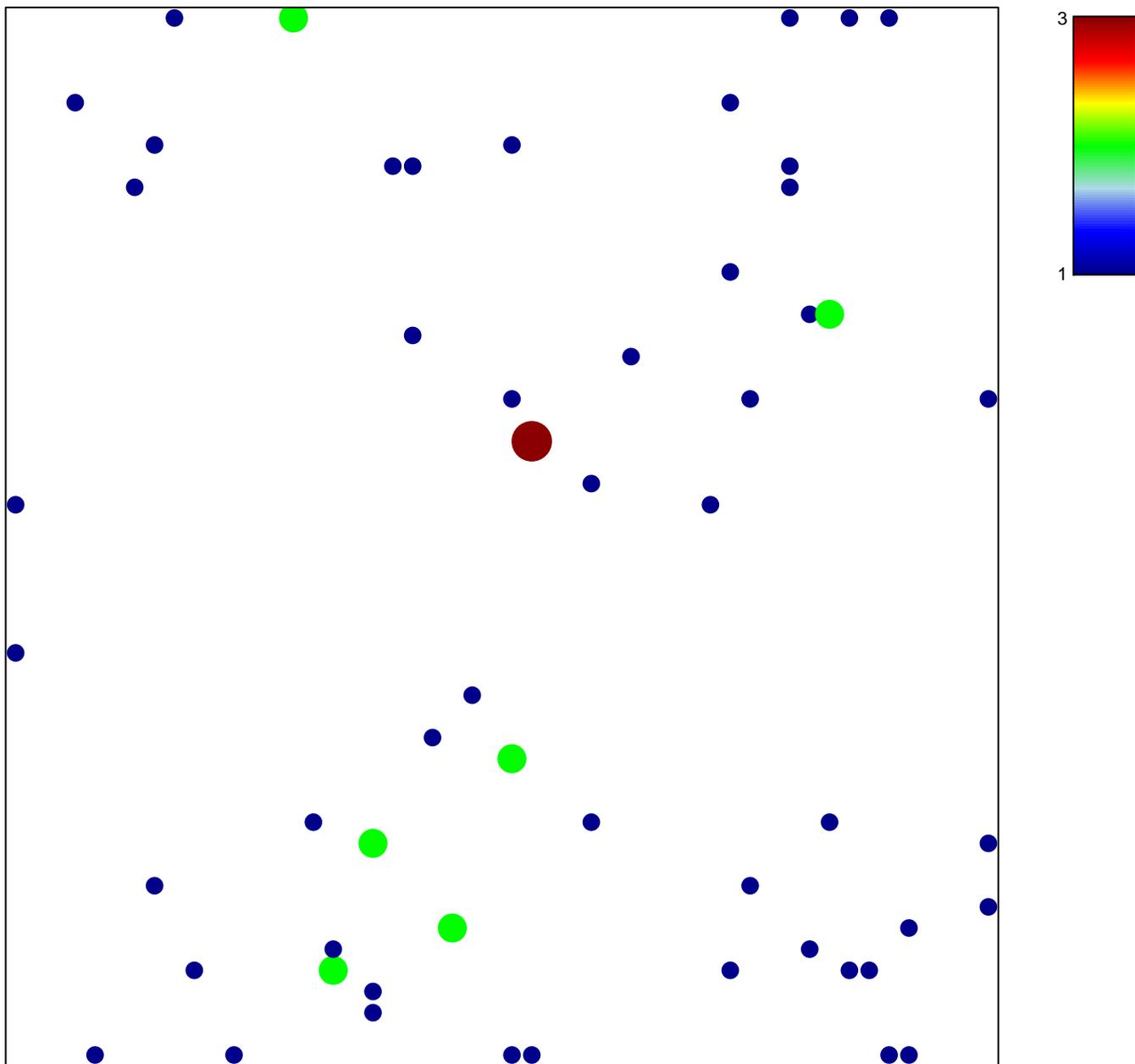
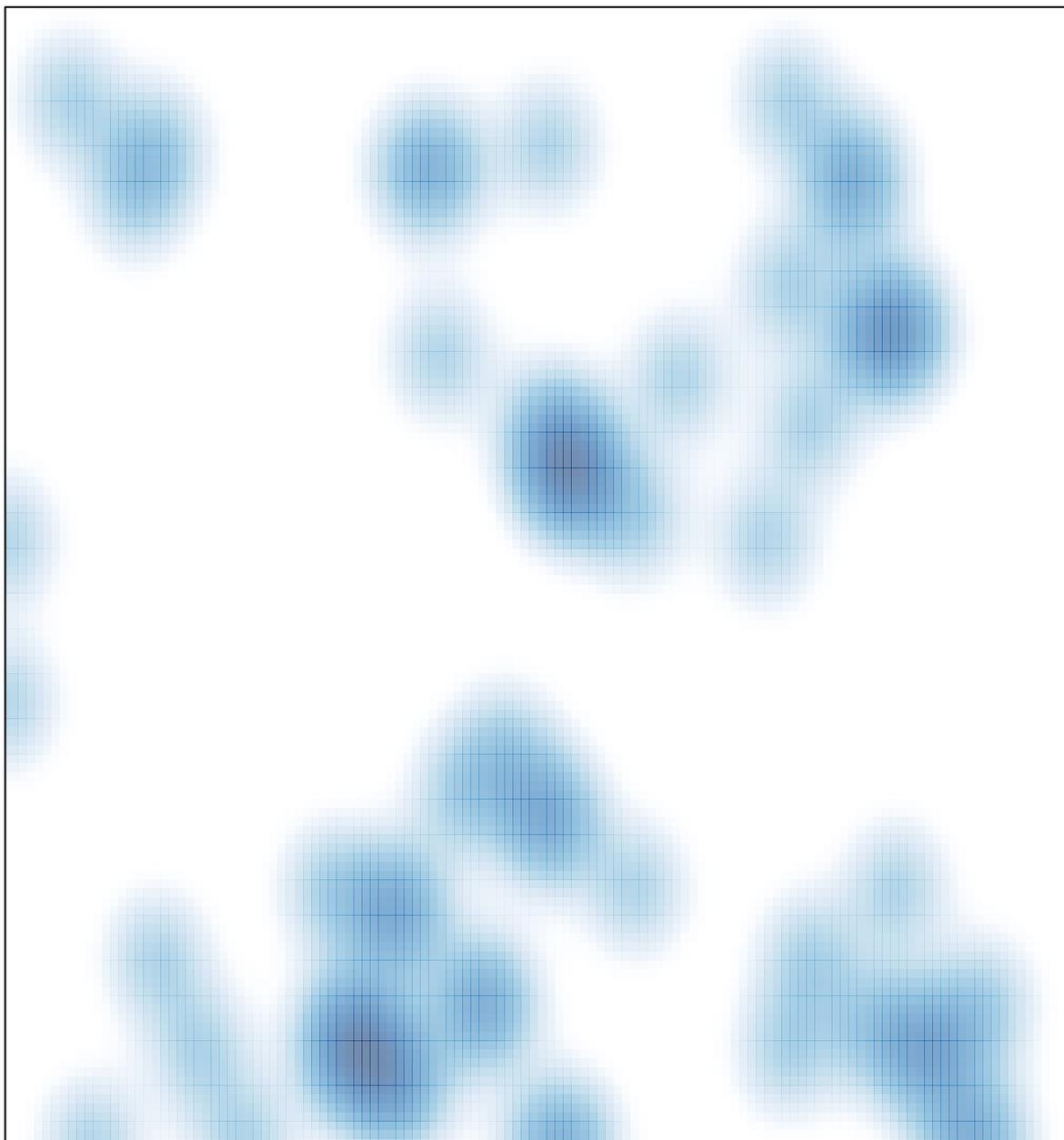


regulatory region sequence-specific DNA binding transcription factor activity involved in



features = 63
chi-square p = 0.8

sequence-specific DNA binding transcription factor a



features = 63 , max = 3