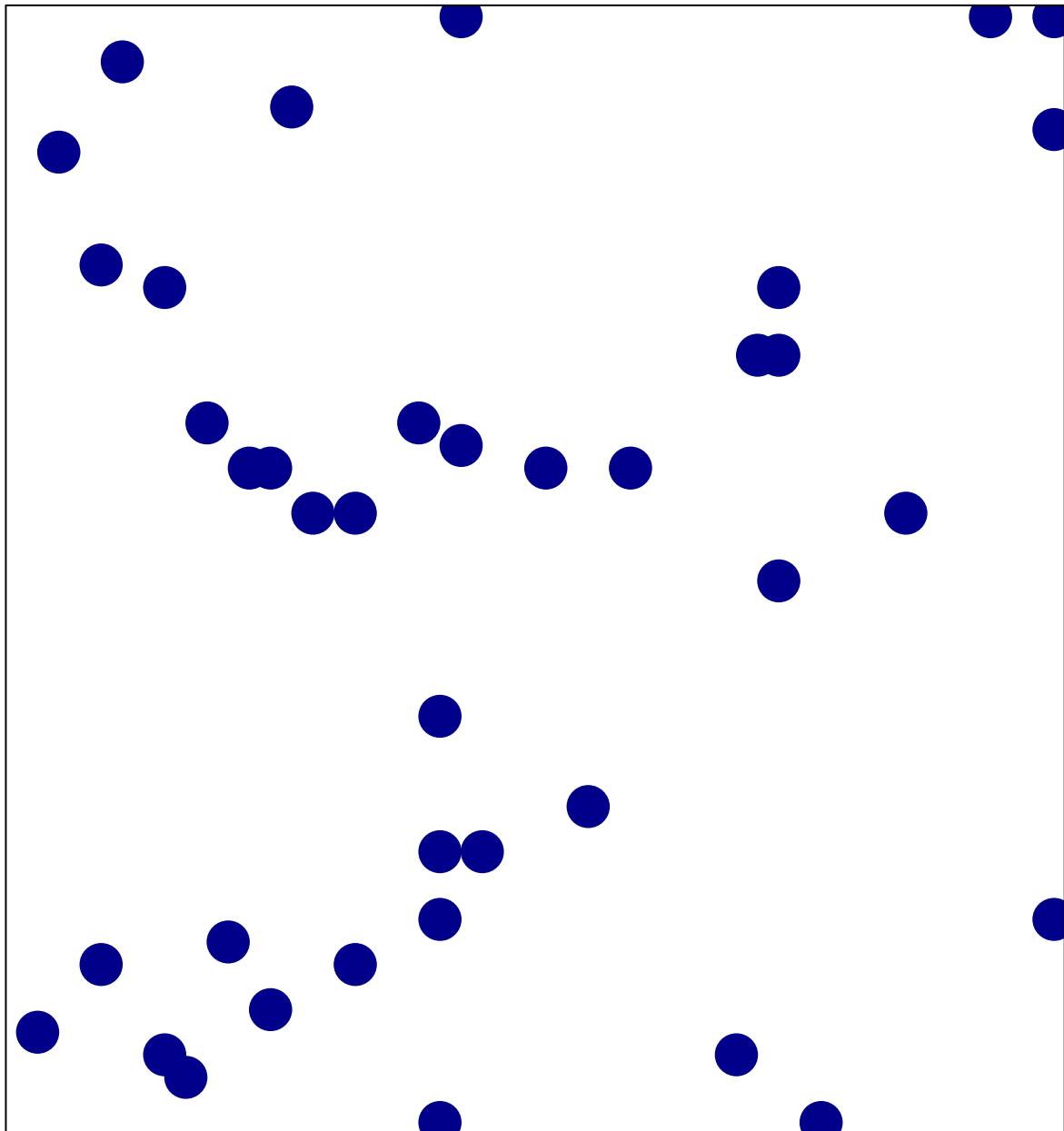
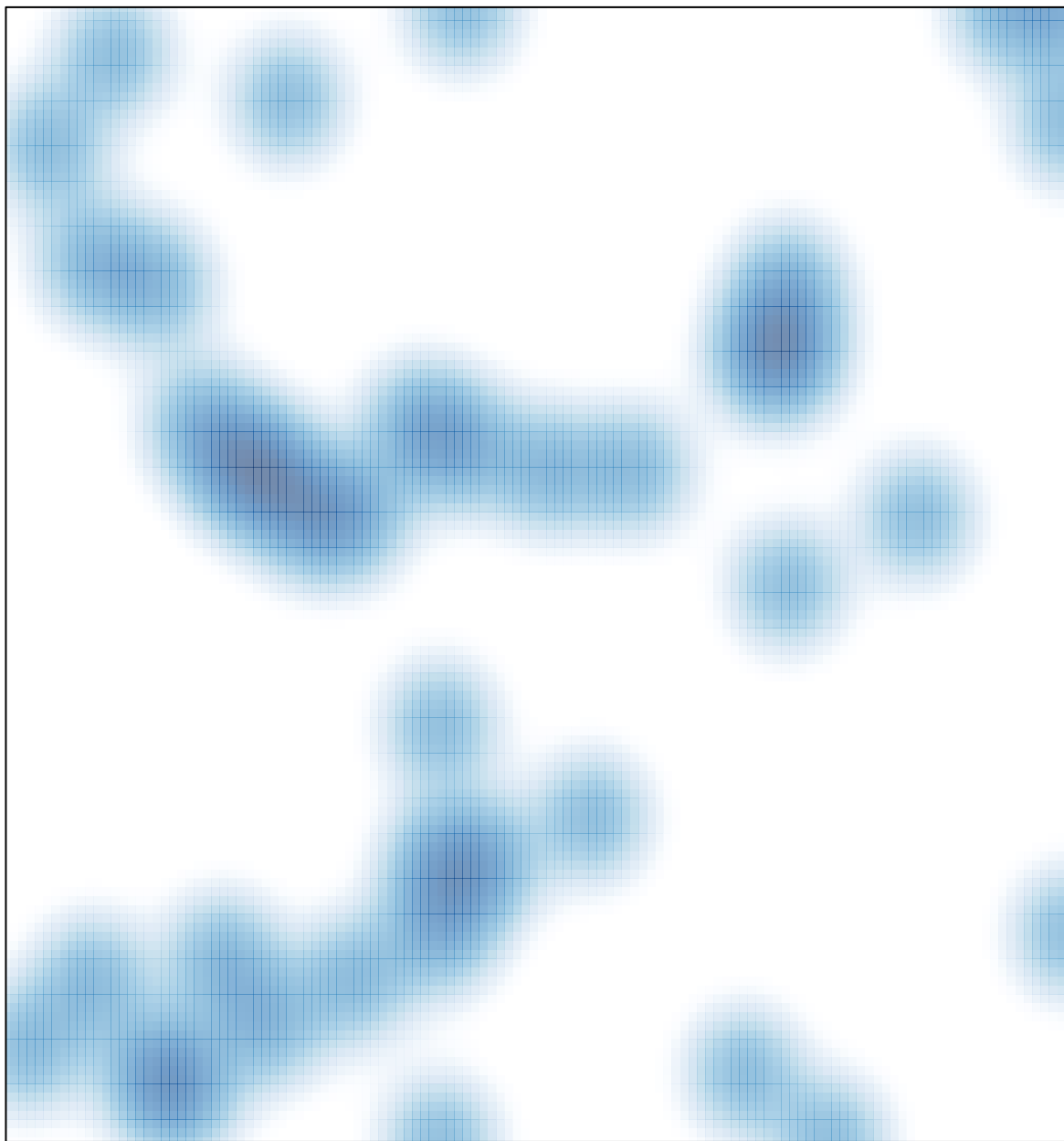


transcription regulatory region sequence-specific DNA binding



features = 39
chi-square p = 0.83

cription regulatory region sequence-specific DNA b



features = 39 , max = 1