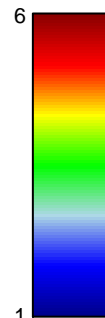
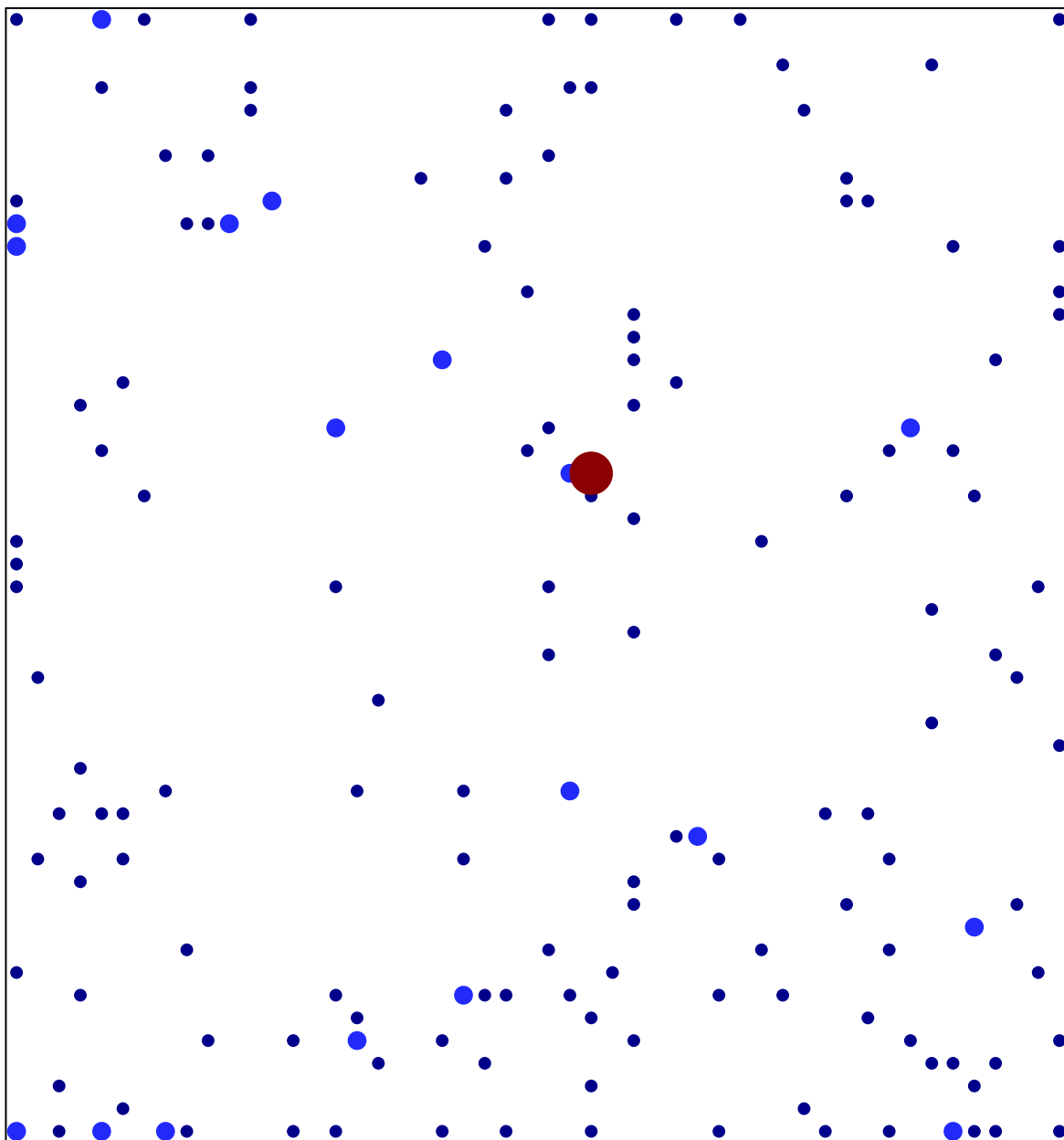
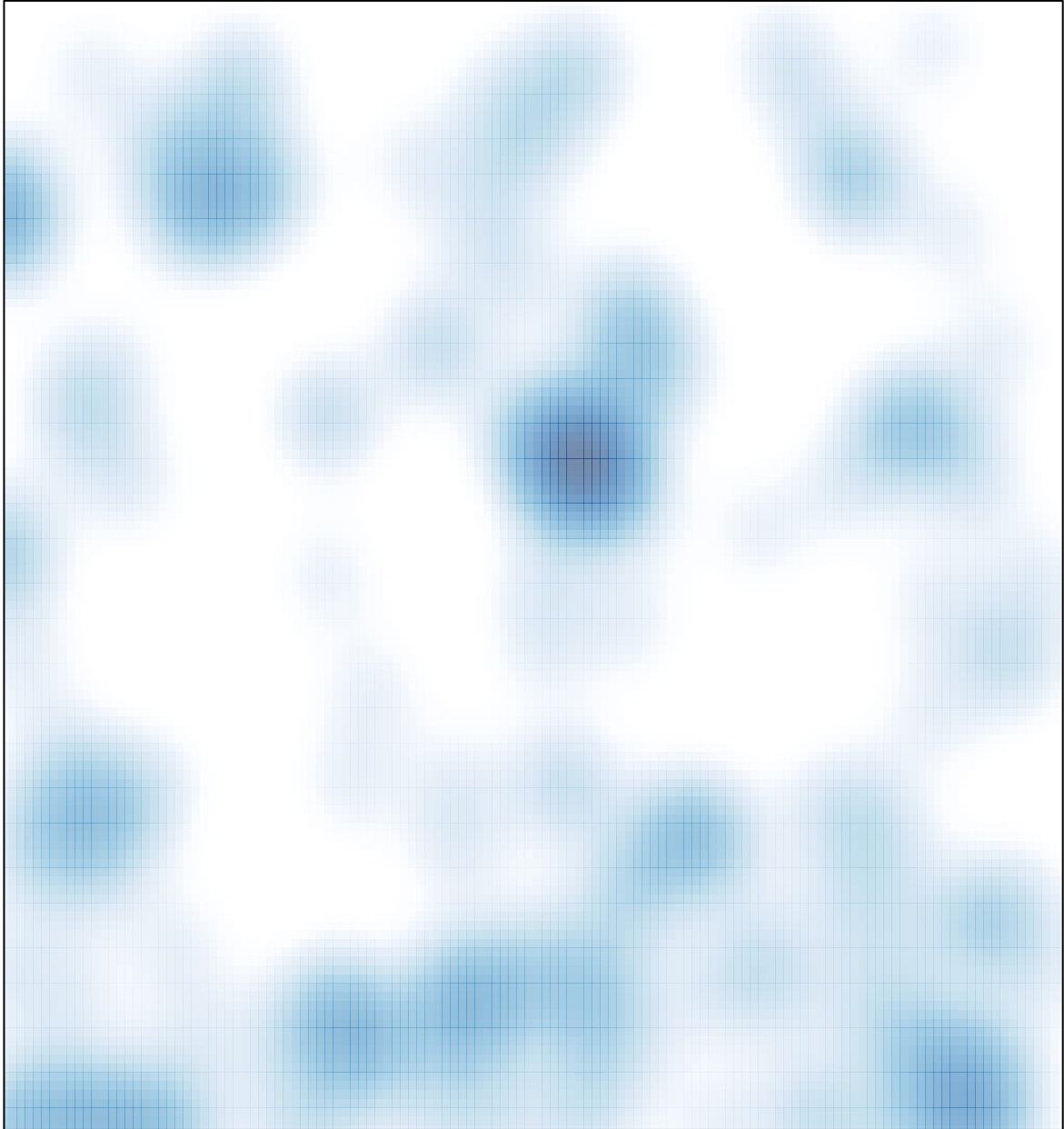


# transcription regulatory region DNA binding



# features = 176  
chi-square p = 0.75

# transcription regulatory region DNA binding



# features = 176 , max = 6