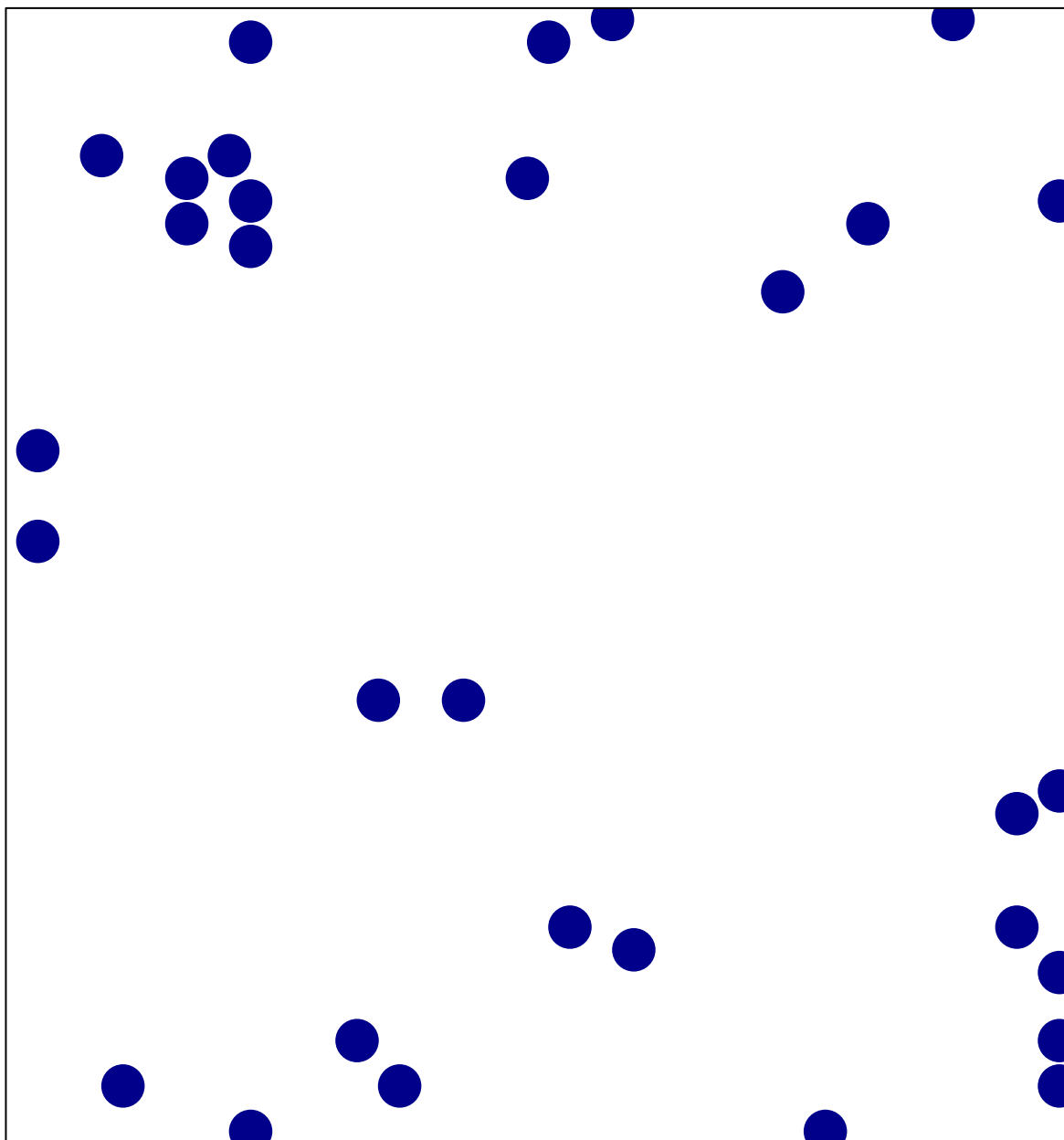
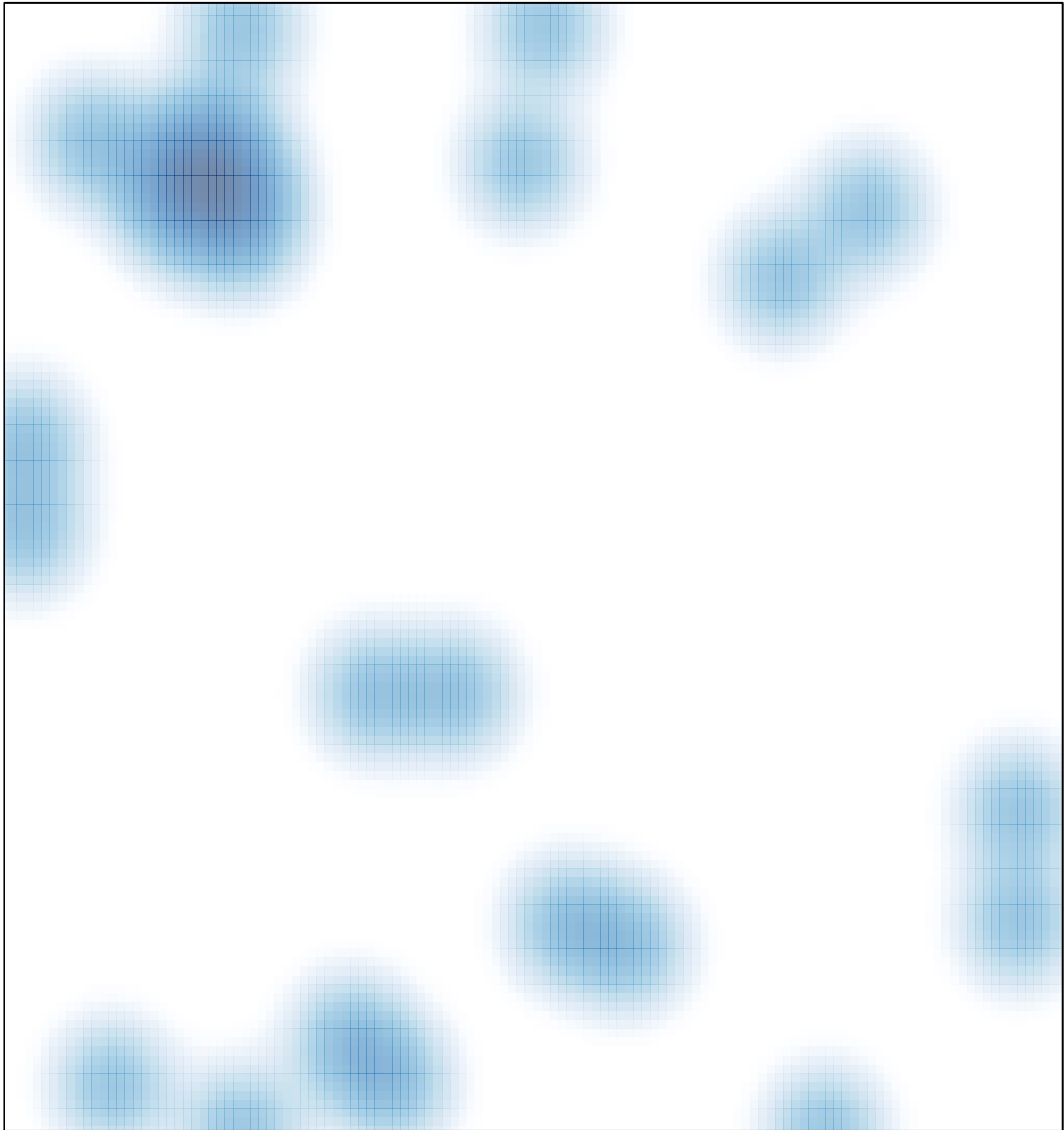


# negative regulation of nucleic acid-templated transcription



# features = 31  
chi-square p = 0.81

# negative regulation of nucleic acid-templated transcrip



# features = 31 , max = 1