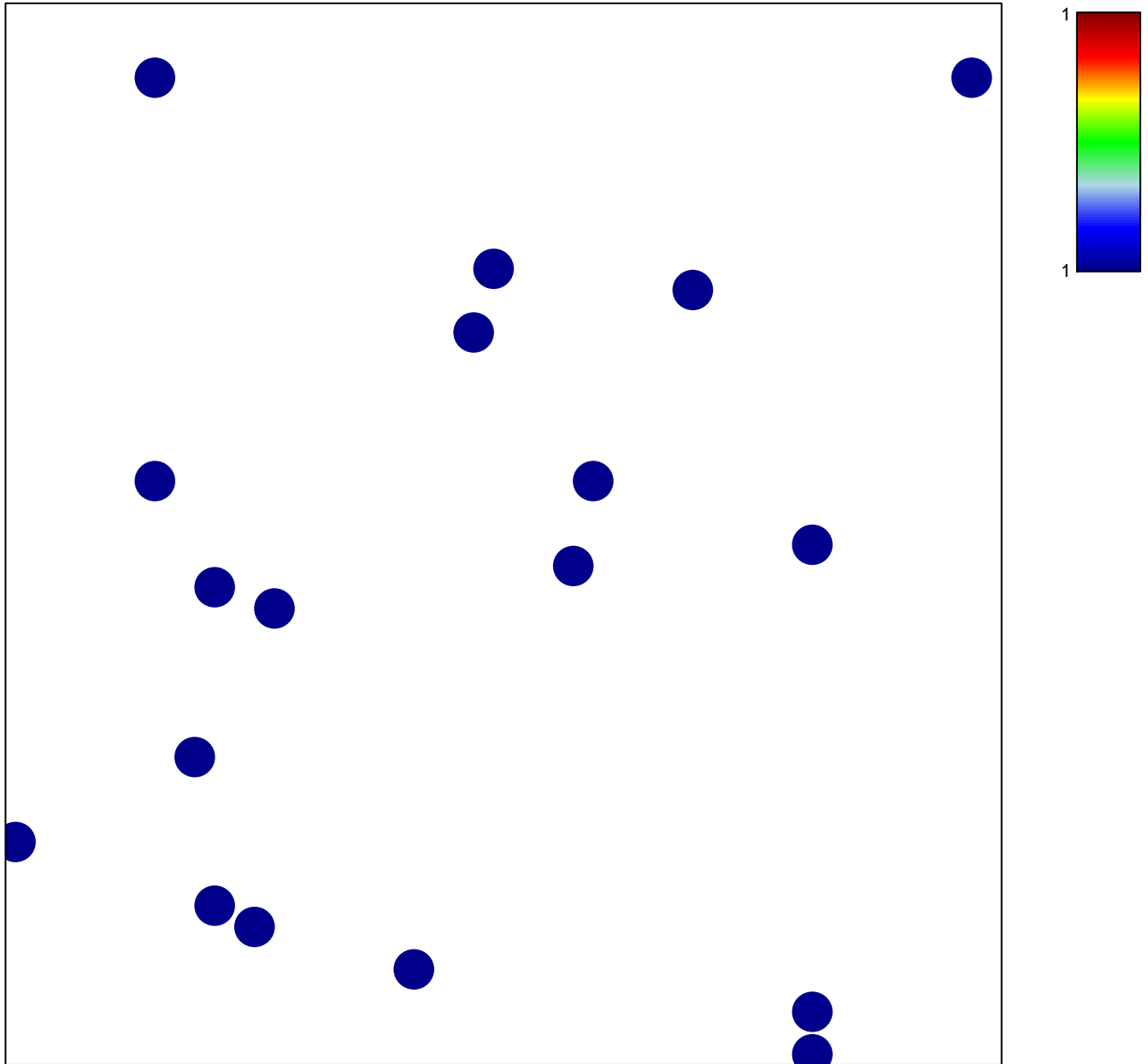
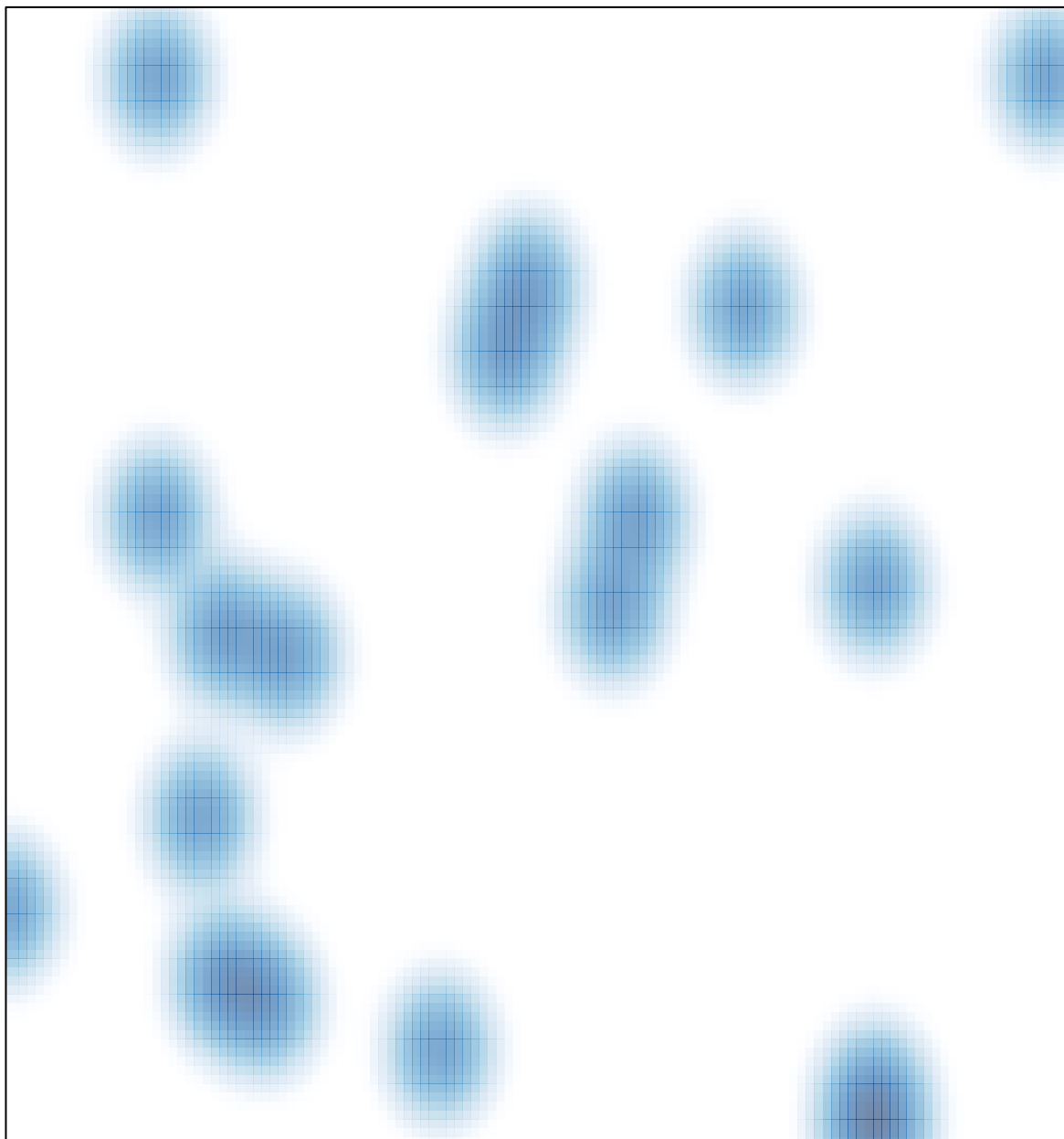


# adenyl nucleotide binding



# features = 18  
chi-square p = 0.84

# adenyl nucleotide binding



# features = 18 , max = 1