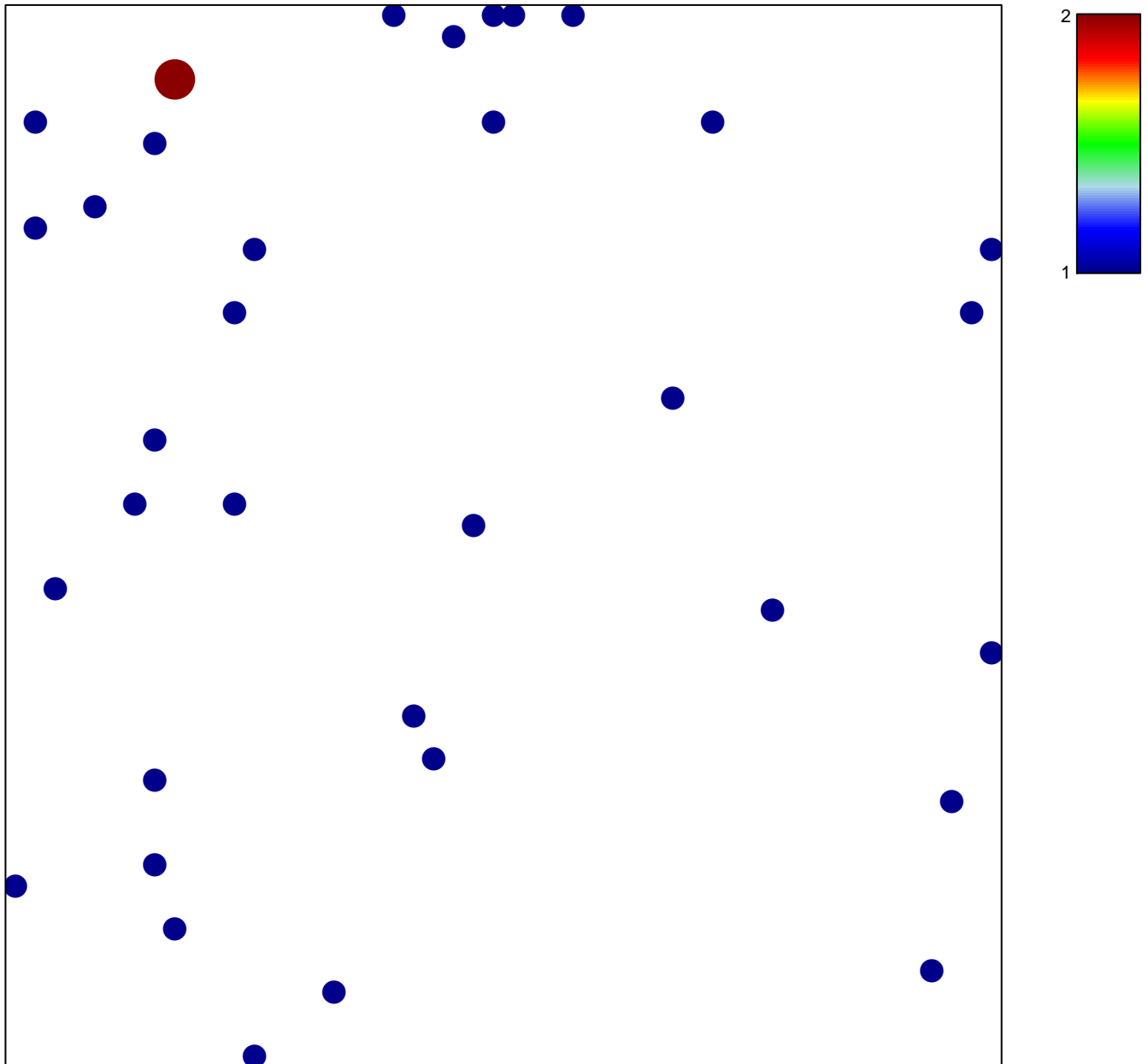
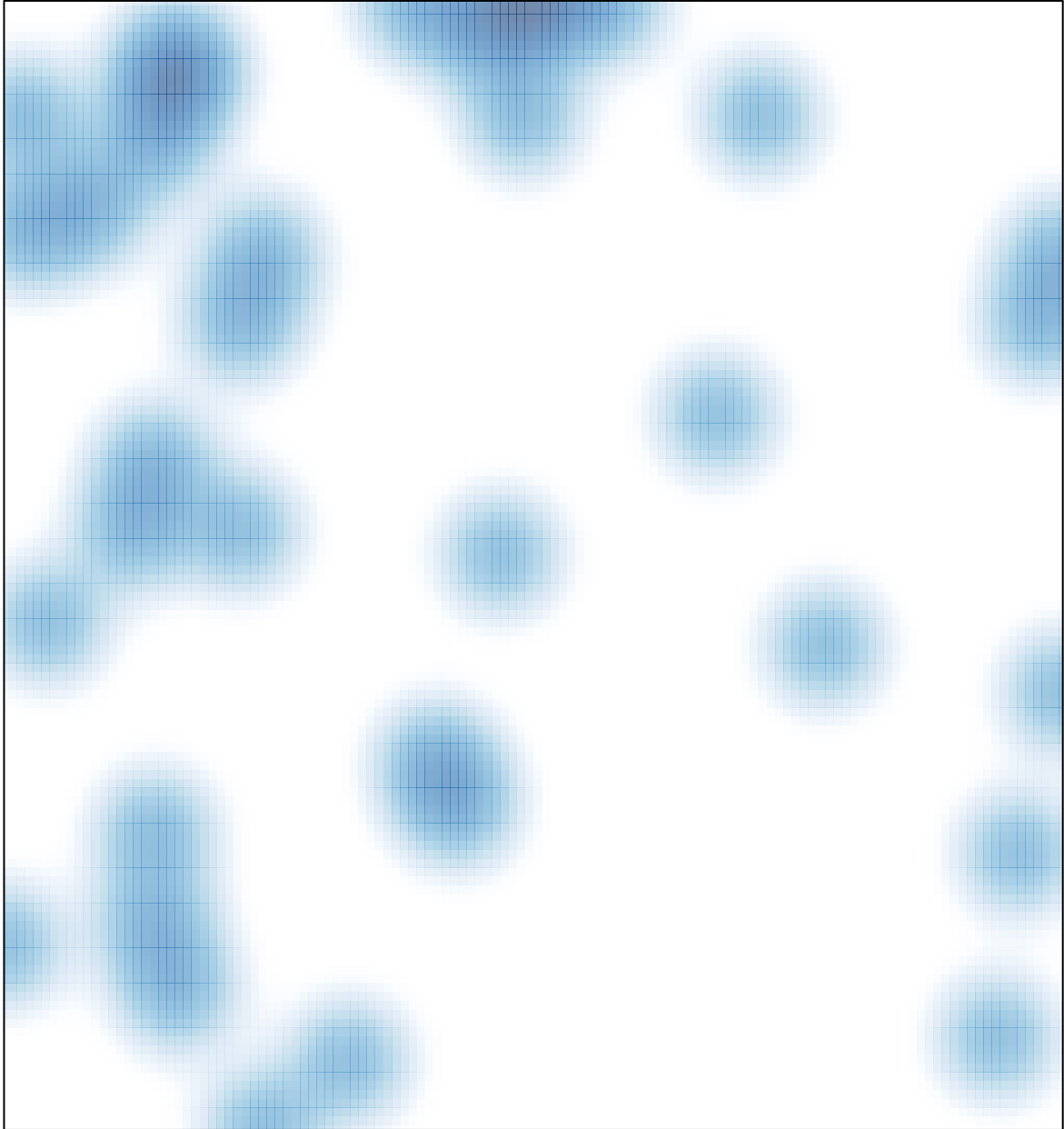


# guanyl nucleotide binding



# features = 35  
chi-square p = 0.84

# guanyl nucleotide binding



# features = 35 , max = 2