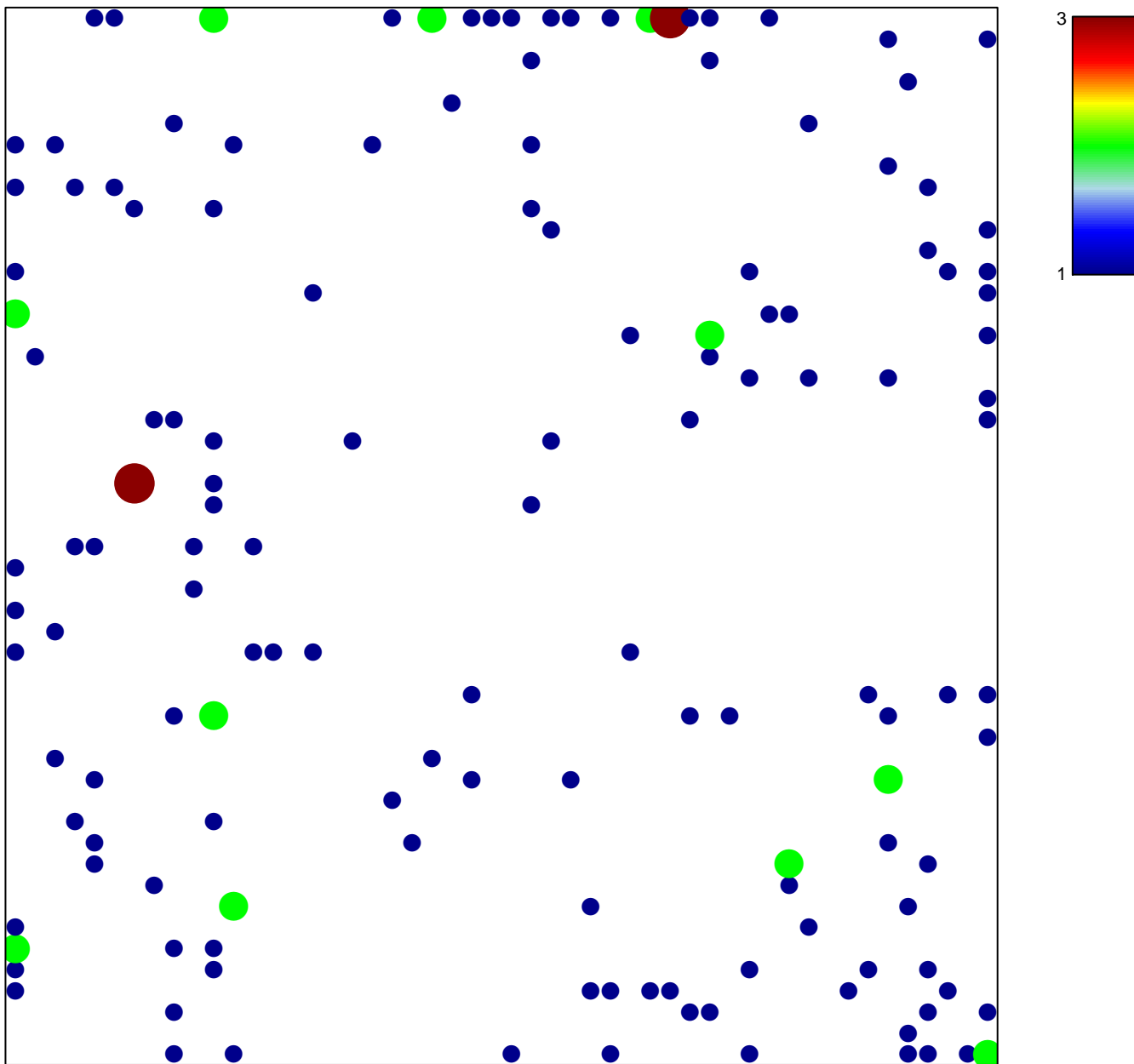
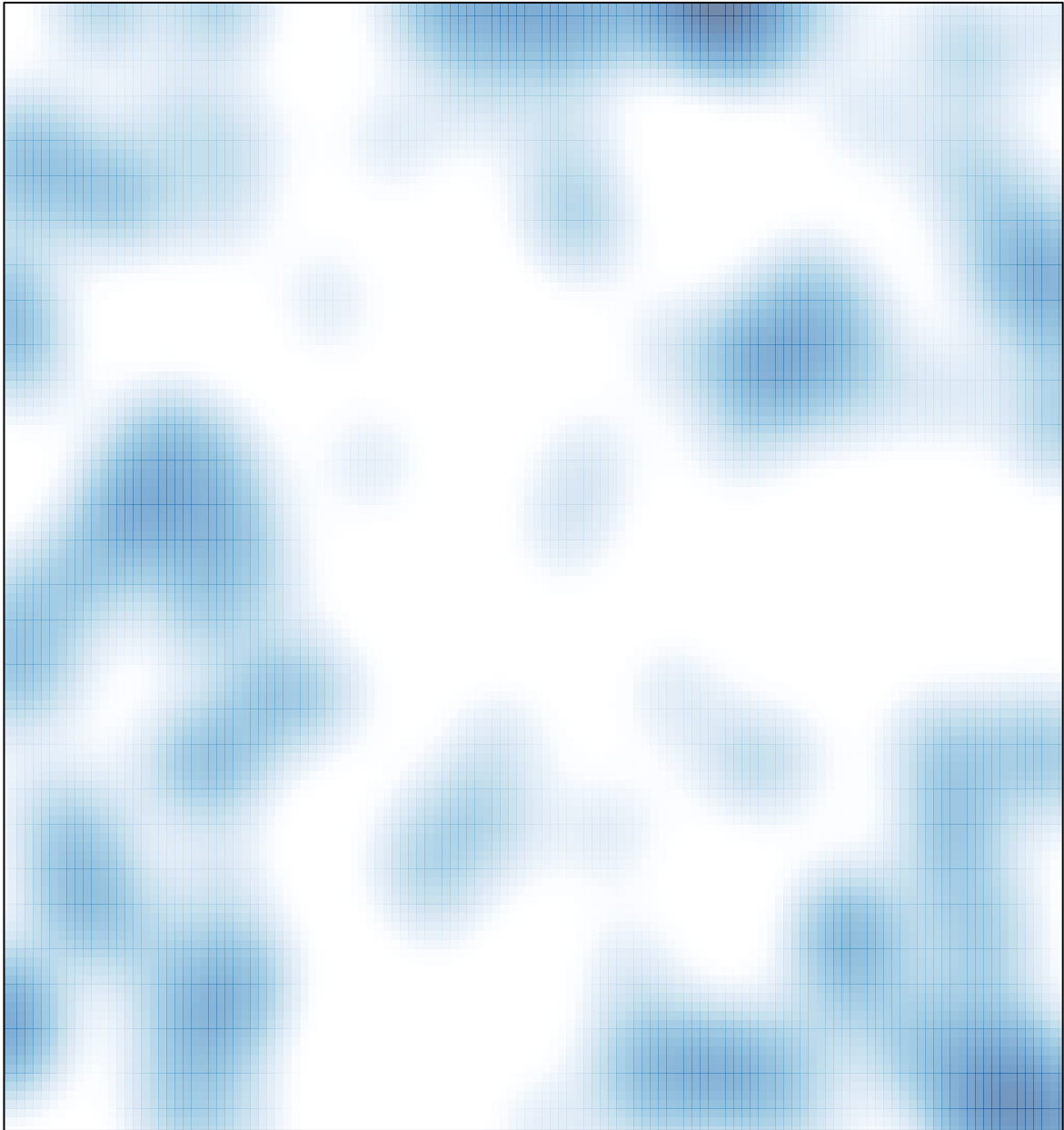


# protein complex binding



# features = 159  
chi-square p = 0.74

# protein complex binding



# features = 159 , max = 3