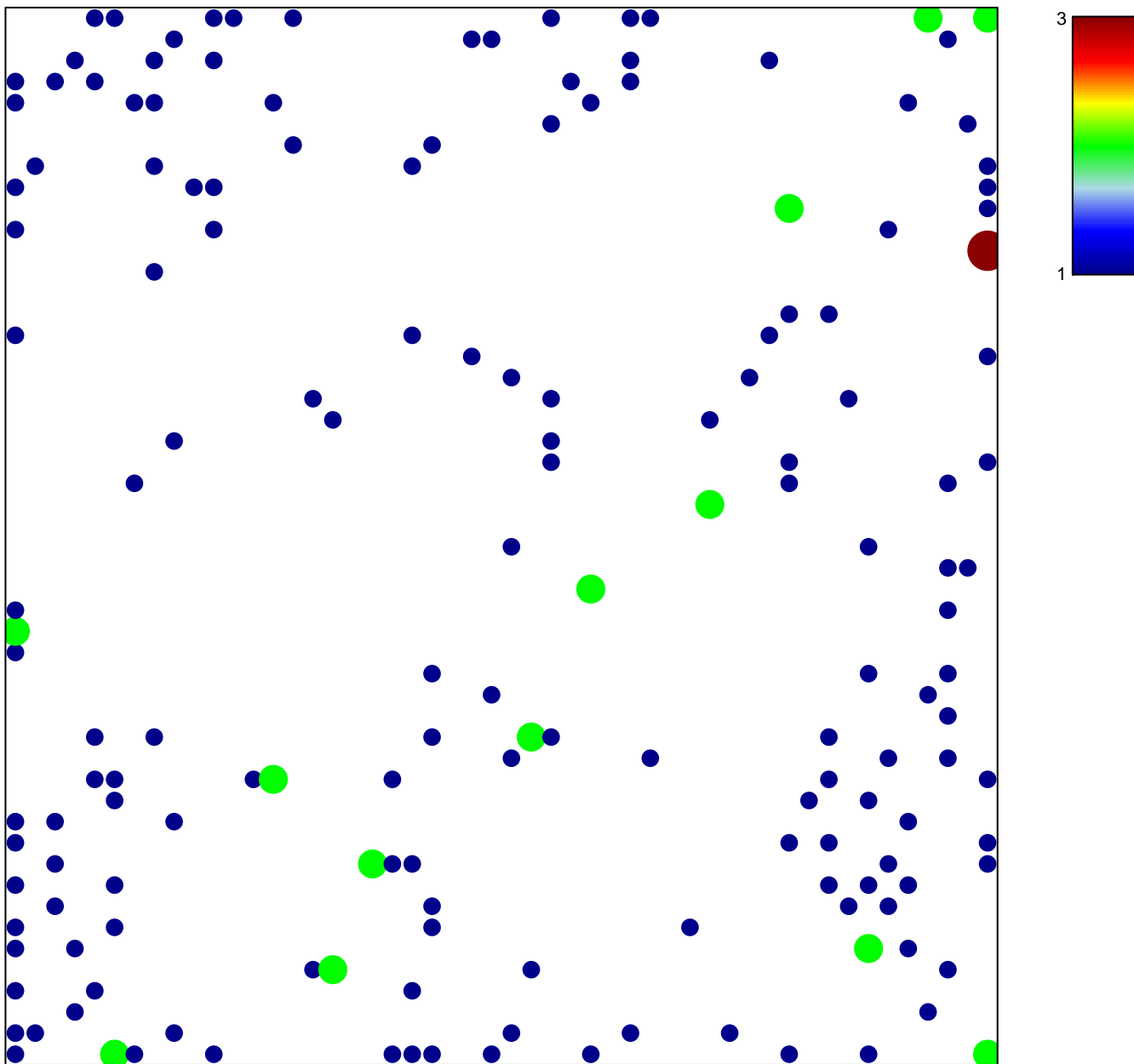
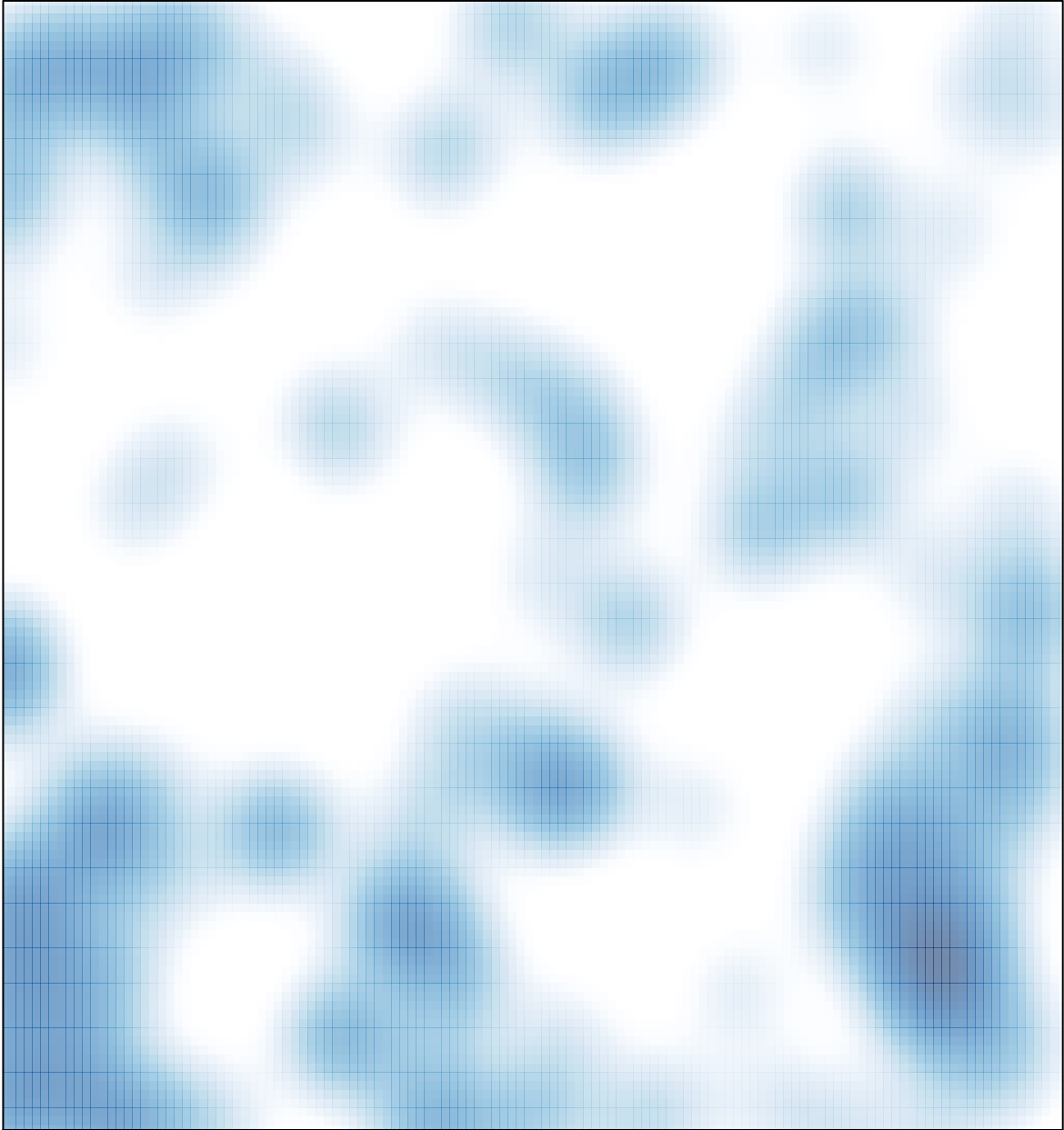


# protein domain specific binding



# features = 180  
chi-square p = 0.76

# protein domain specific binding



# features = 180 , max = 3