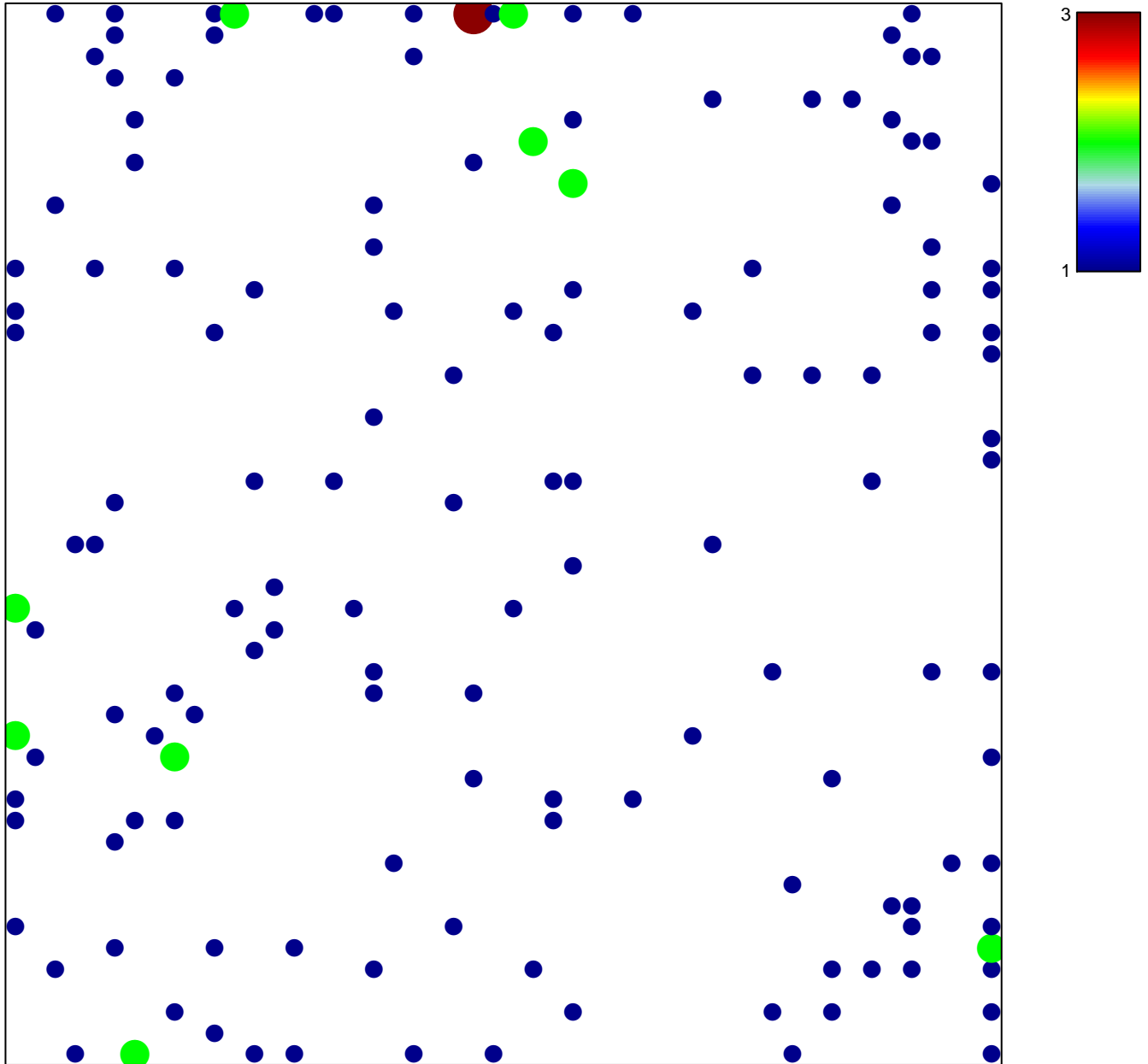
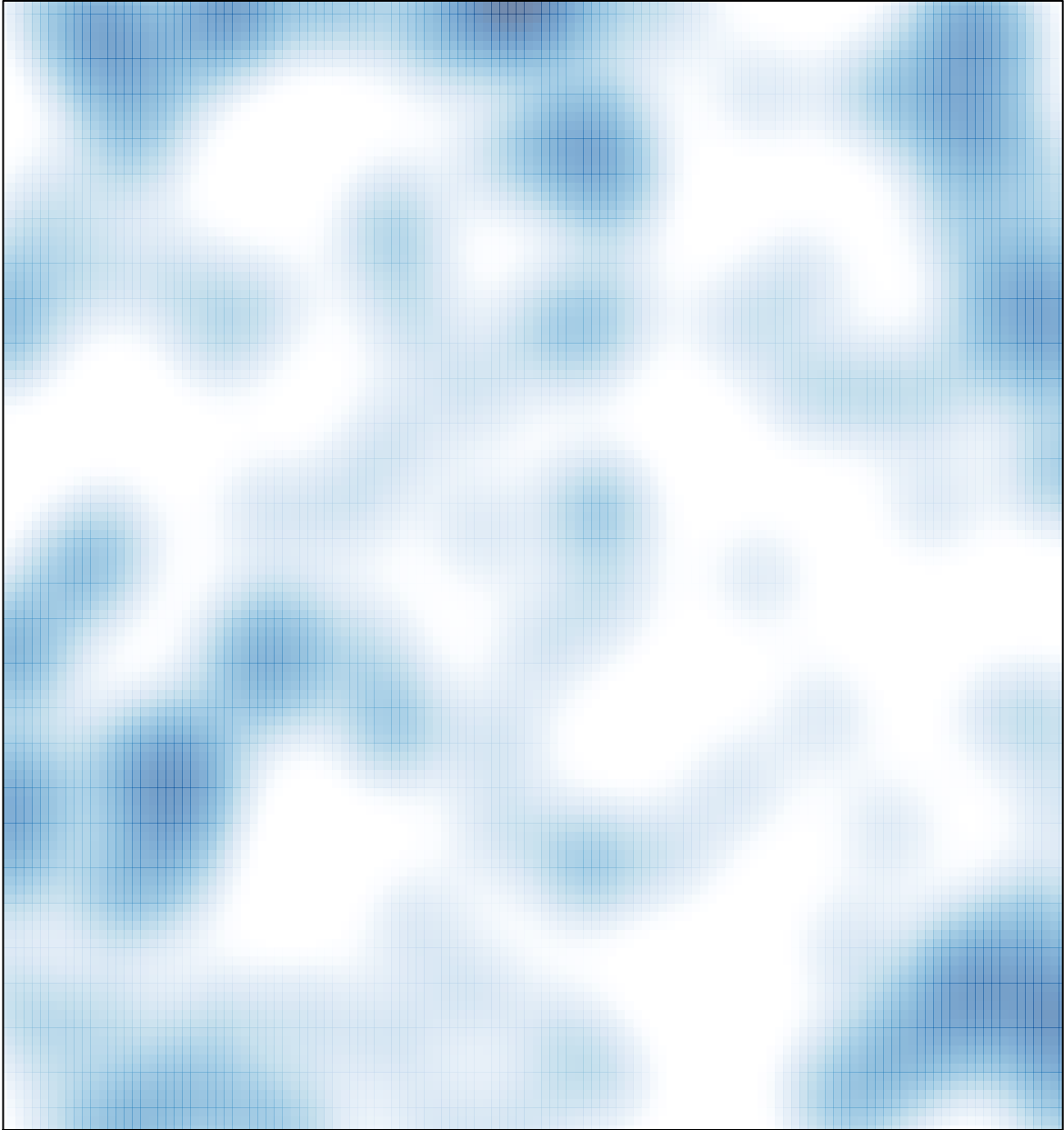


# protein C-terminus binding



# features = 156  
chi-square p = 0.75

# protein C-terminus binding



# features = 156 , max = 3