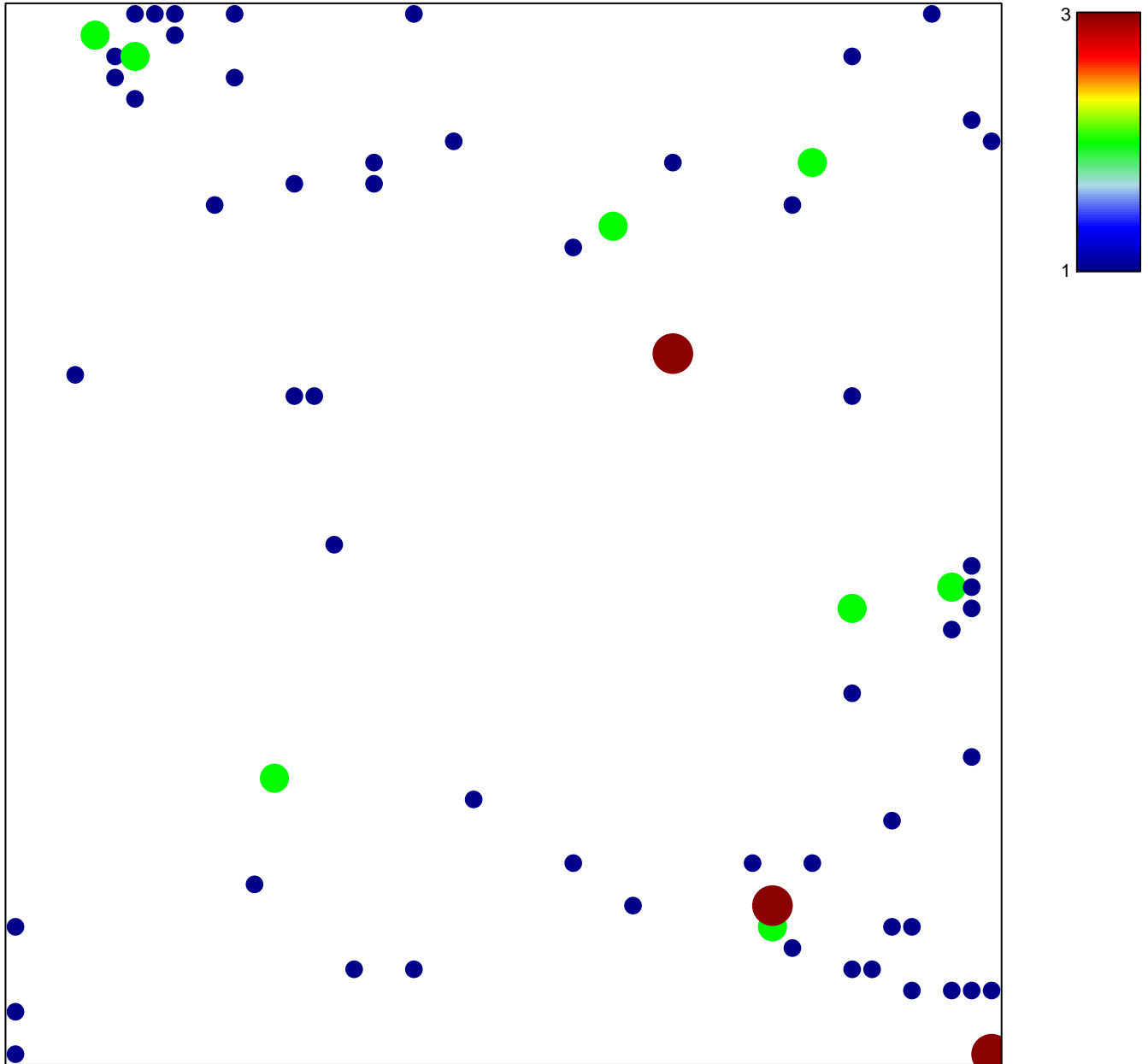
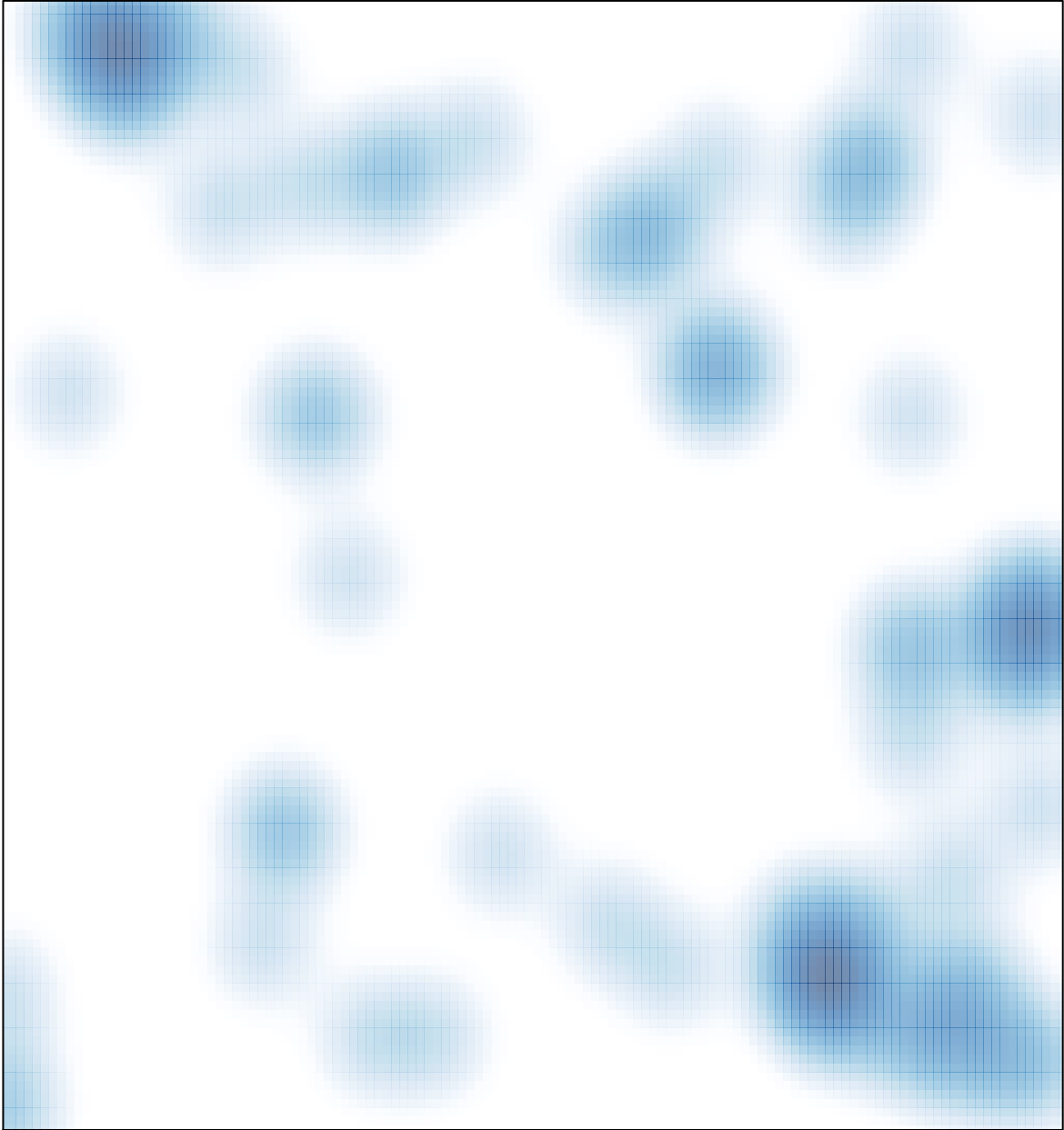


# nucleosome



# features = 79  
chi-square p = 0.8

# nucleosome



# features = 79 , max = 3