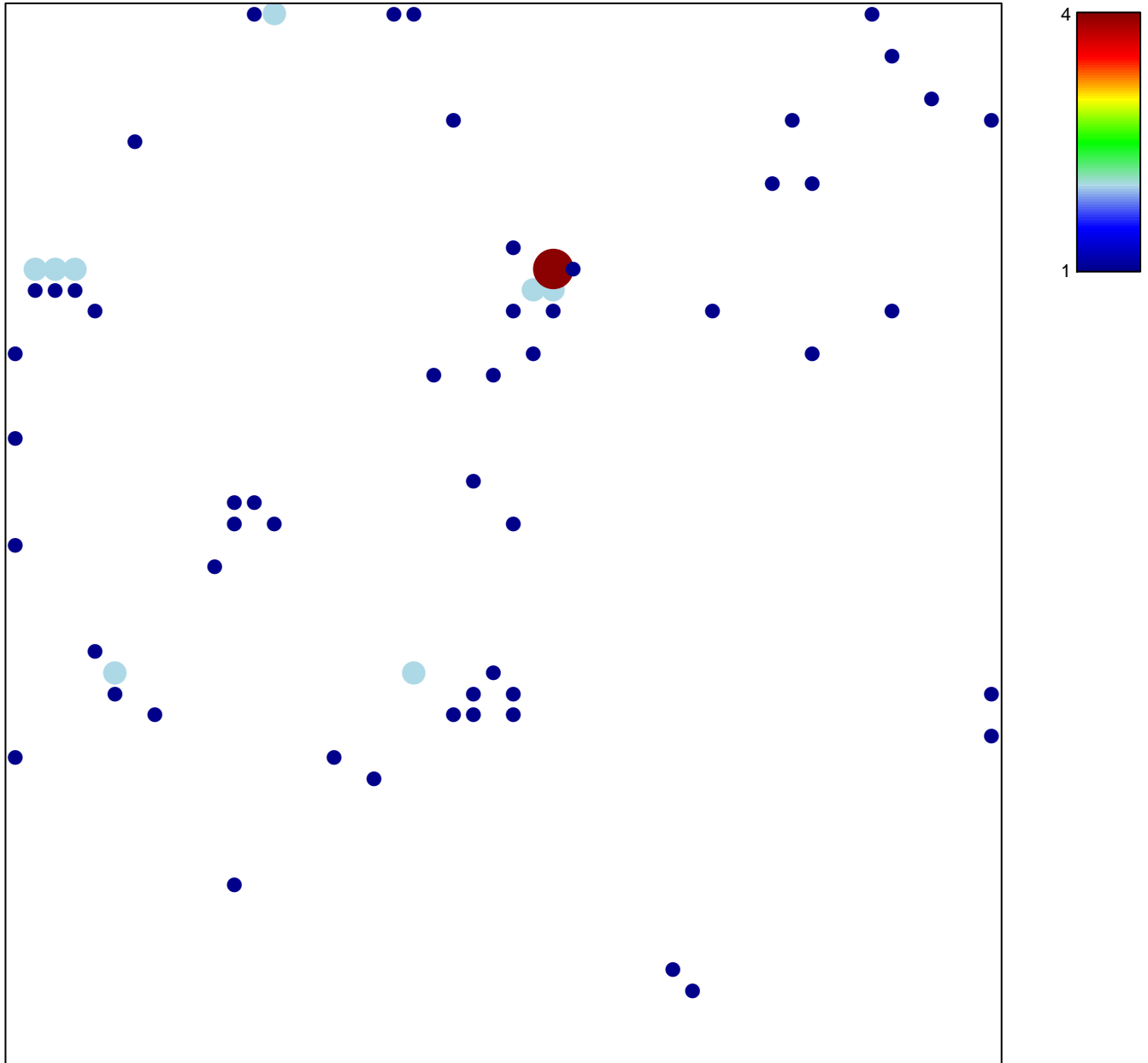
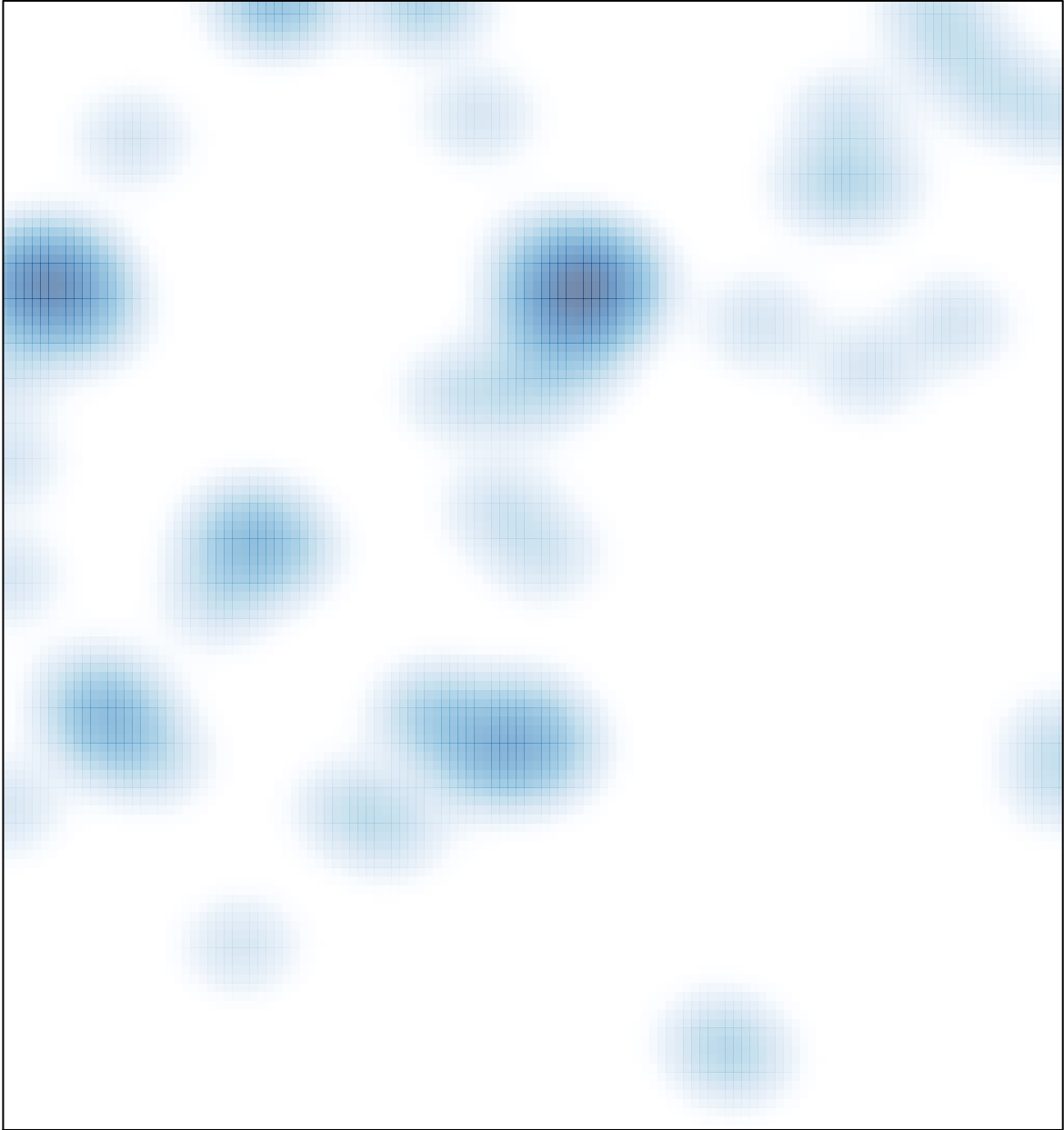


# nucleosome



# features = 73  
chi-square p = 0.76

# nucleosome



# features = 73 , max = 4