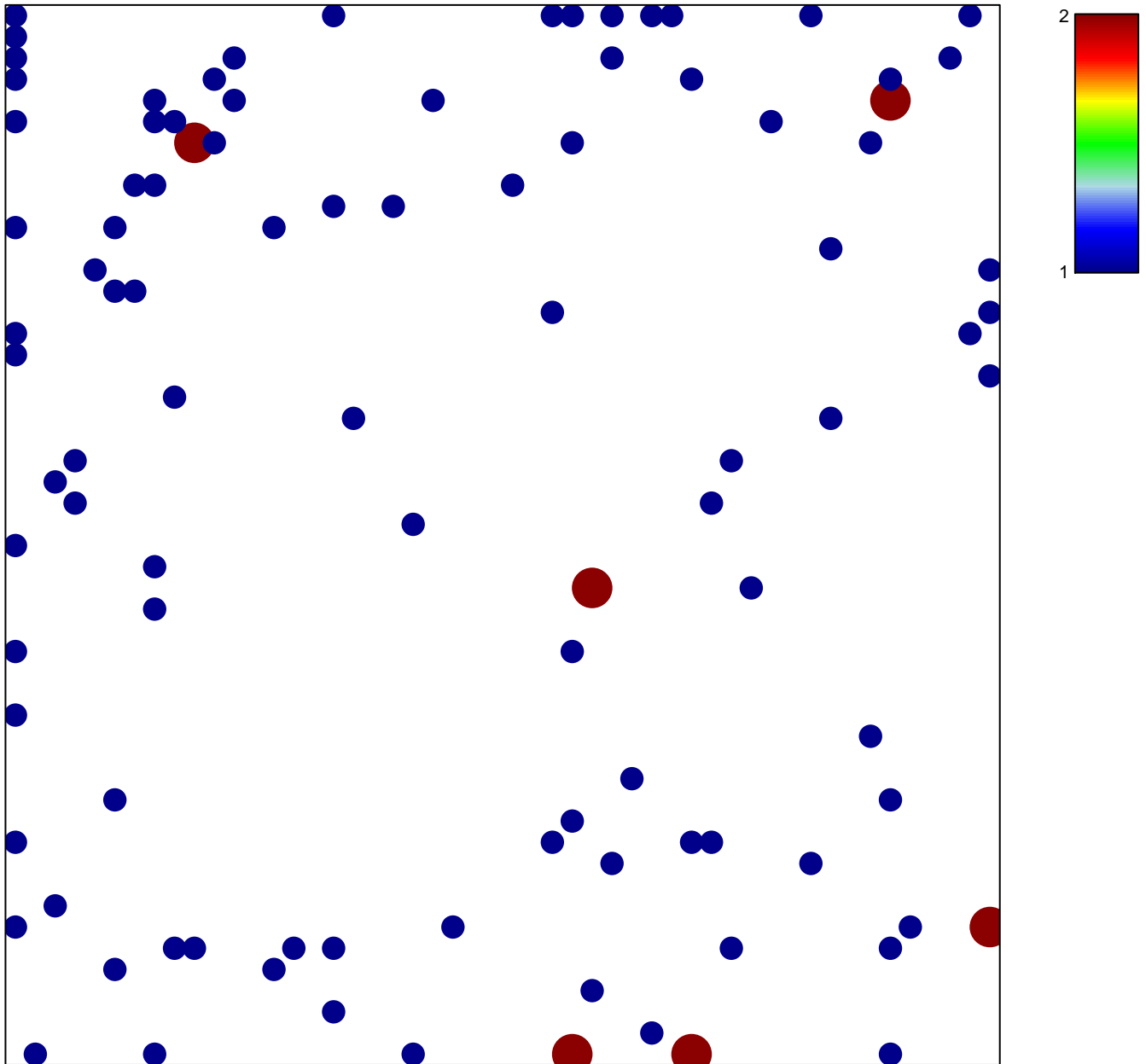
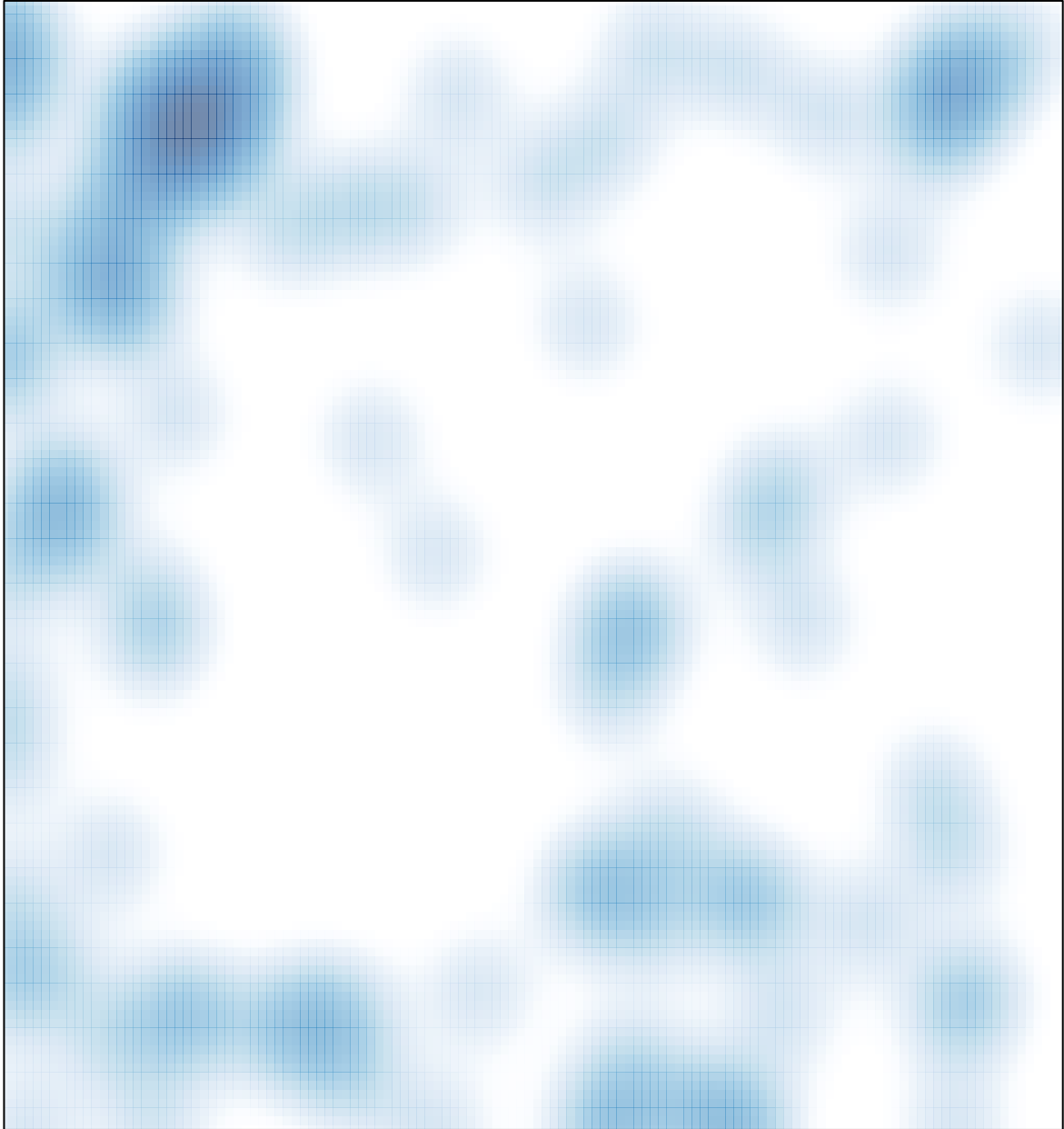


# ribonucleoprotein complex



# features = 105  
chi-square p = 0.77

# ribonucleoprotein complex



# features = 105 , max = 2