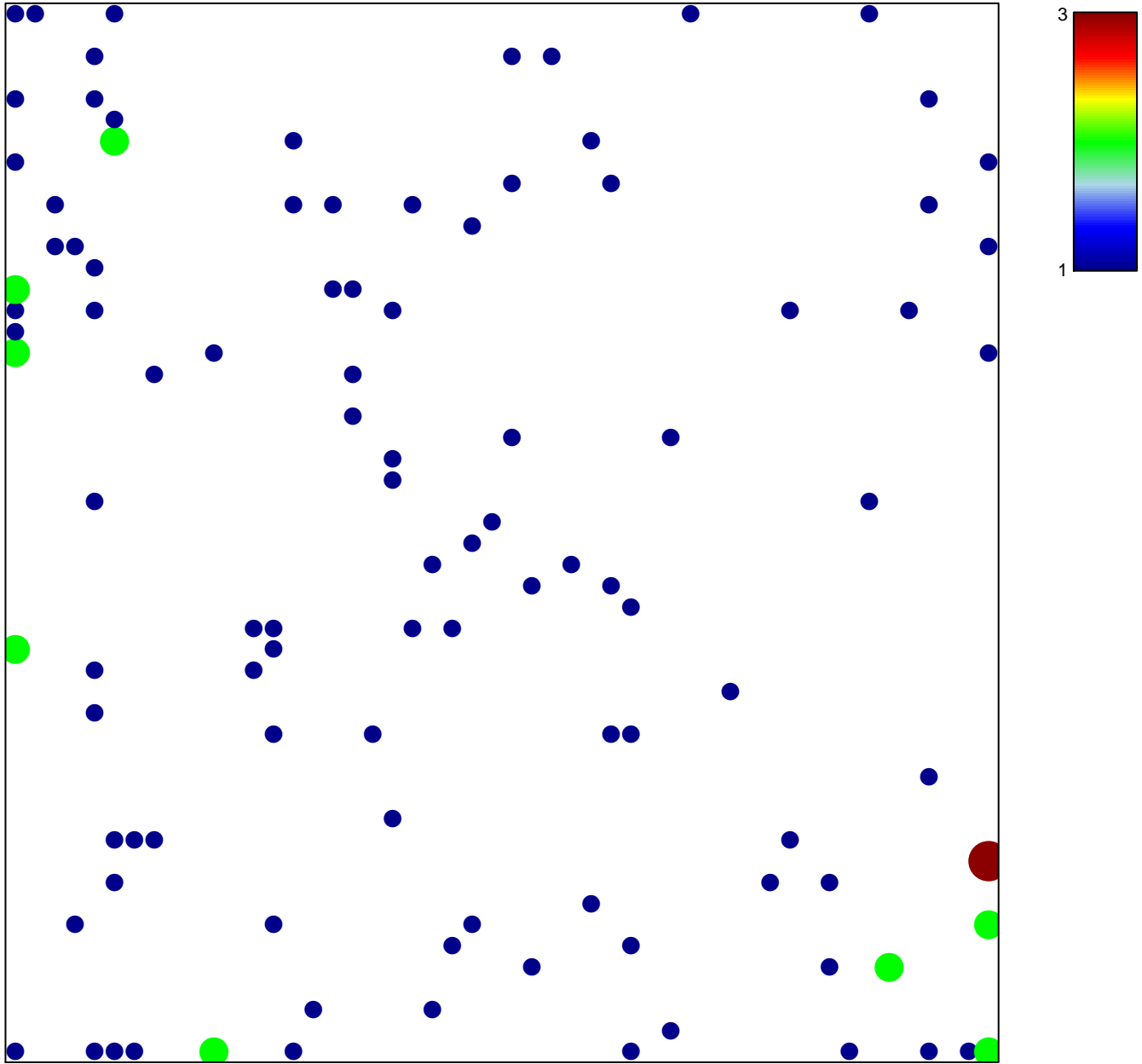
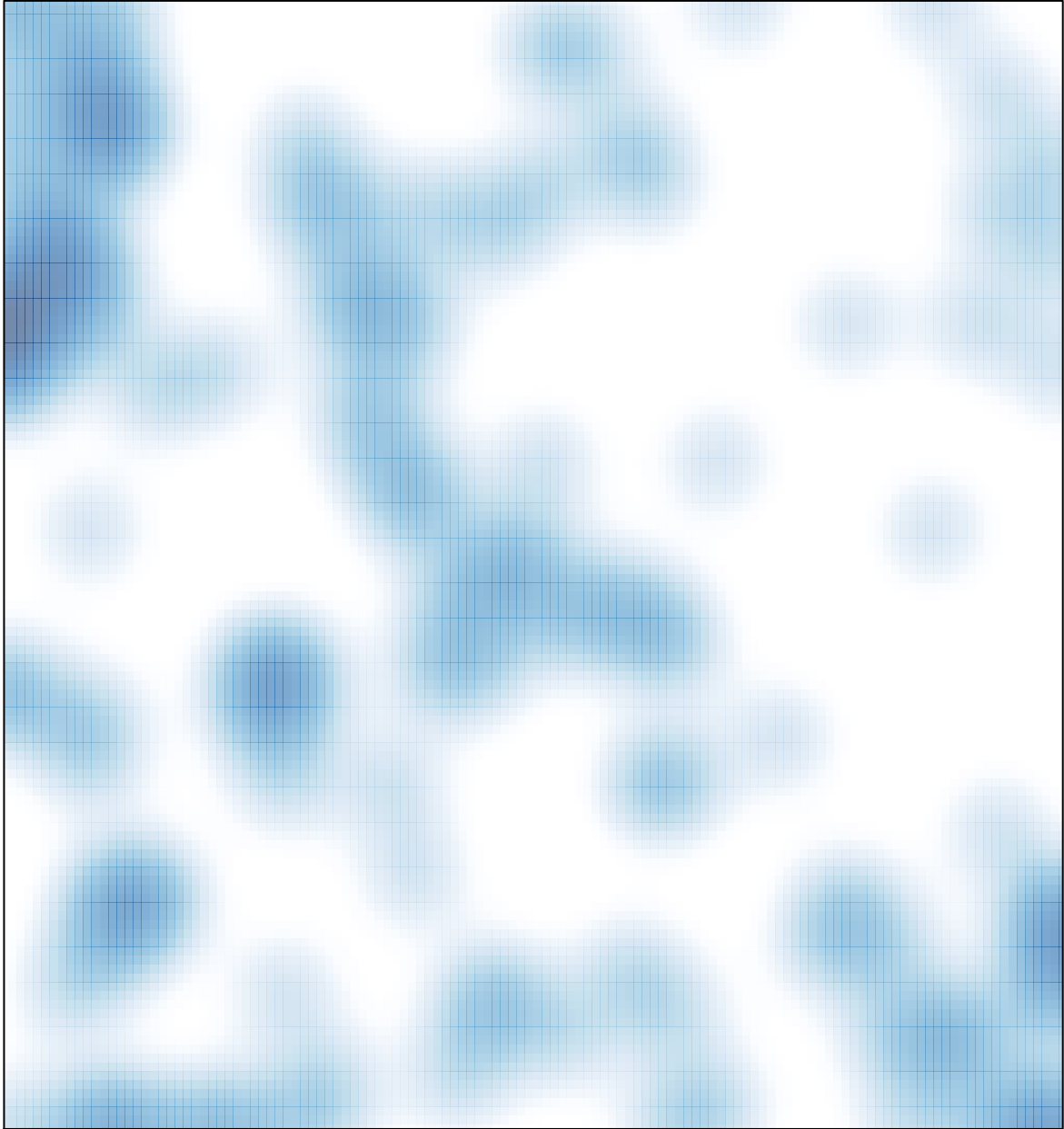


# lipid binding



# features = 115  
chi-square p = 0.77

# lipid binding



# features = 115 , max = 3