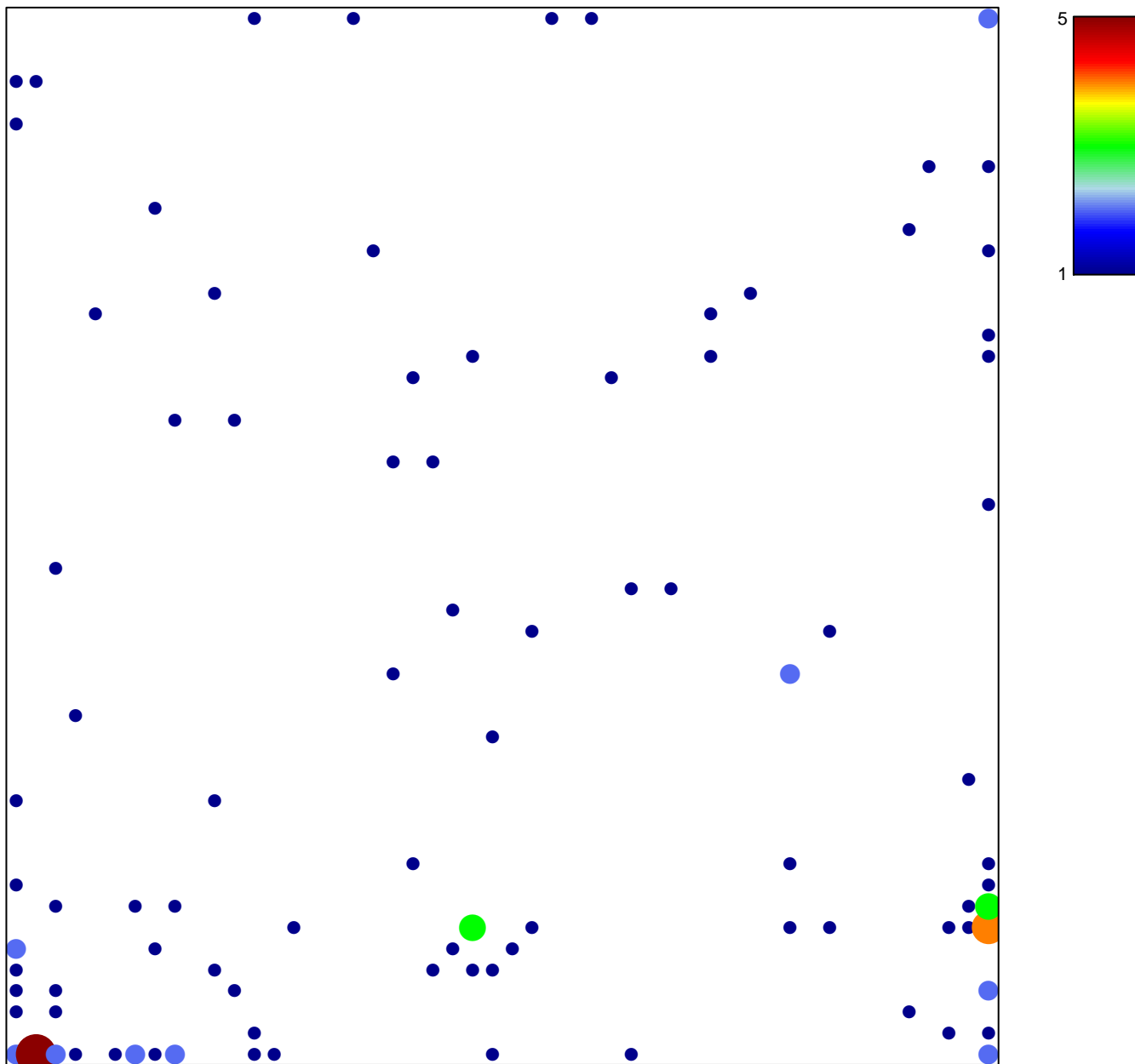
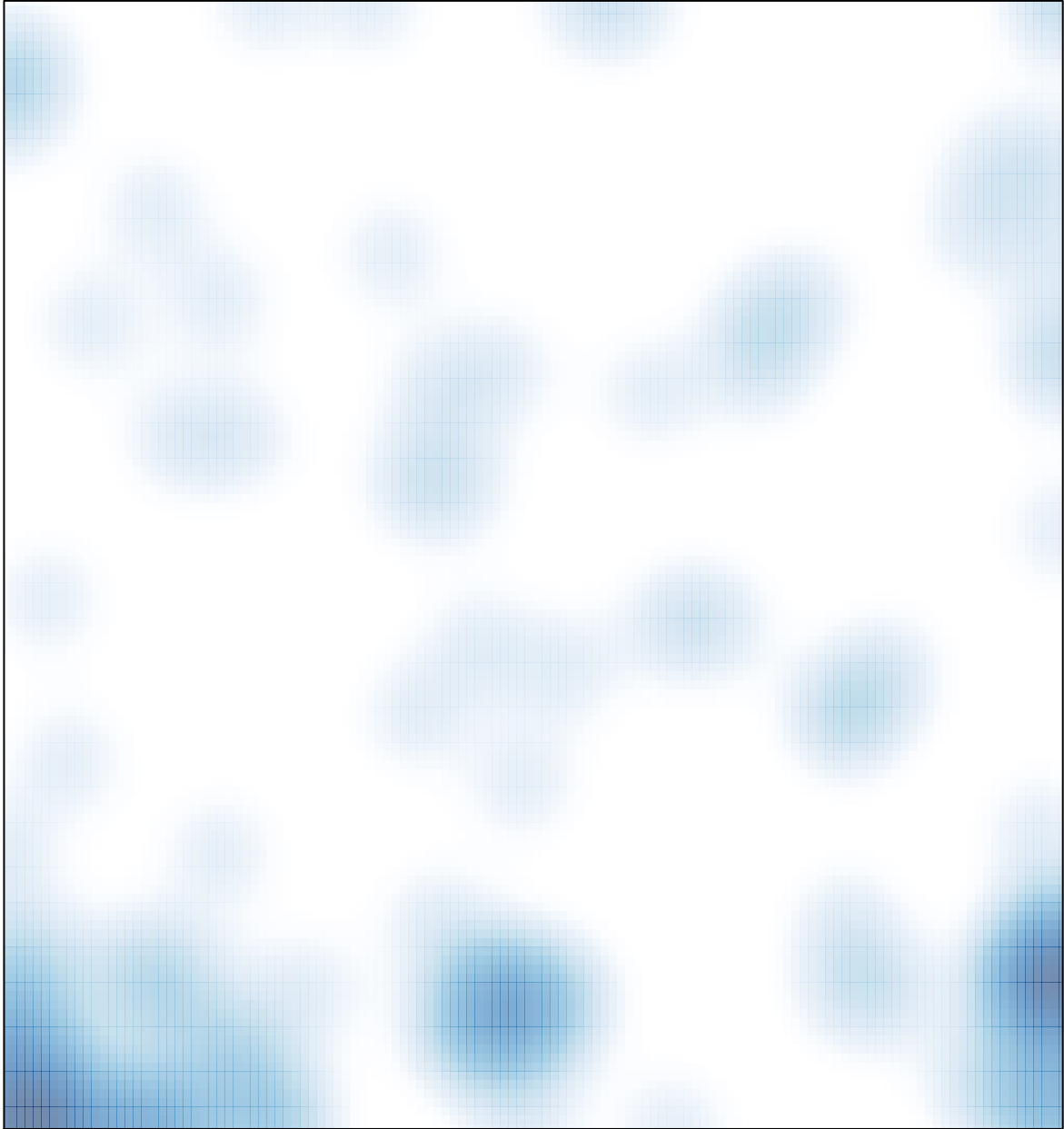


# heparin binding



# features = 112  
chi-square p = 0.64

# heparin binding



# features = 112 , max = 5