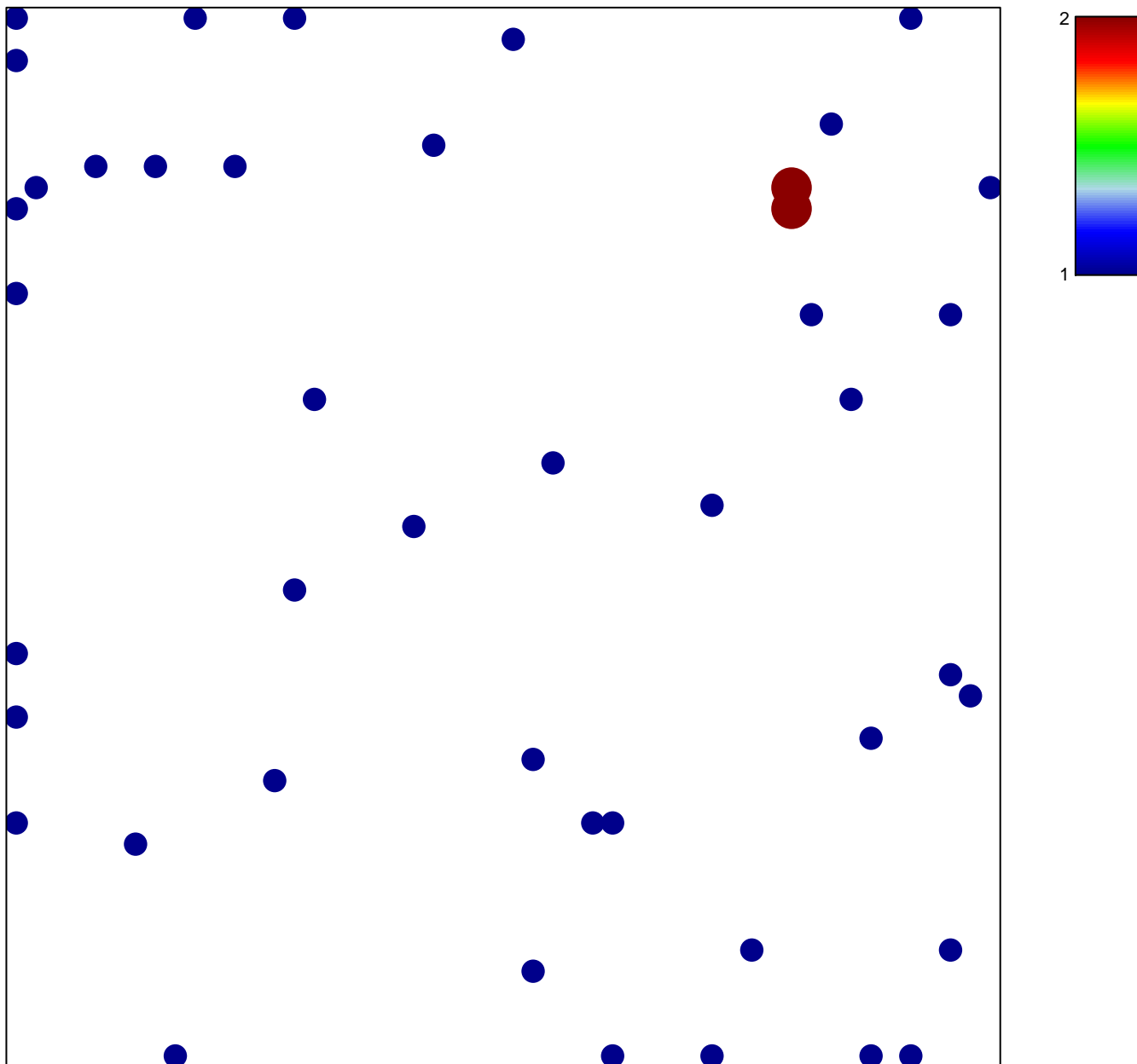
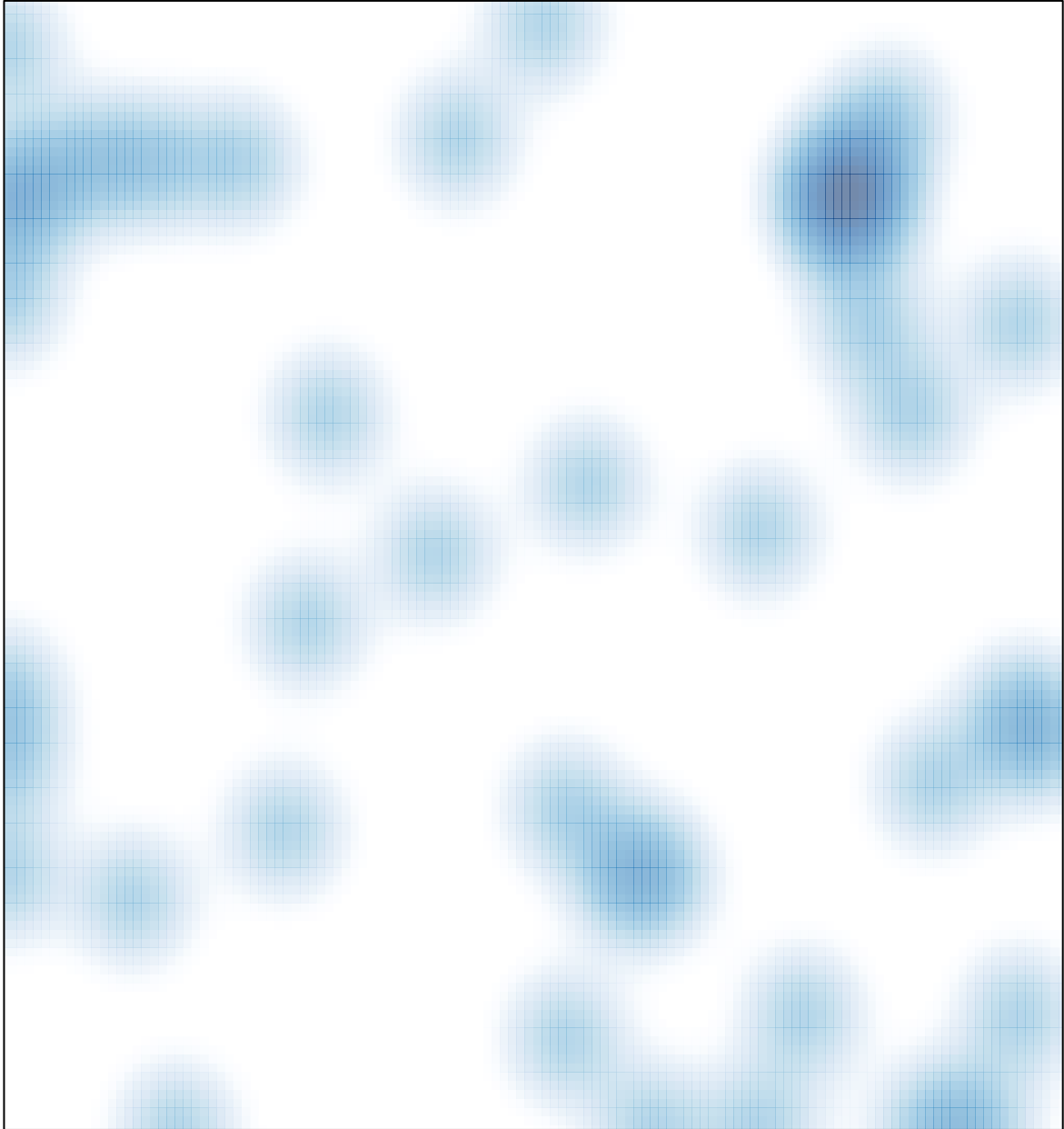


# manganese ion binding



# features = 46  
chi-square p = 0.8

# manganese ion binding



# features = 46 , max = 2