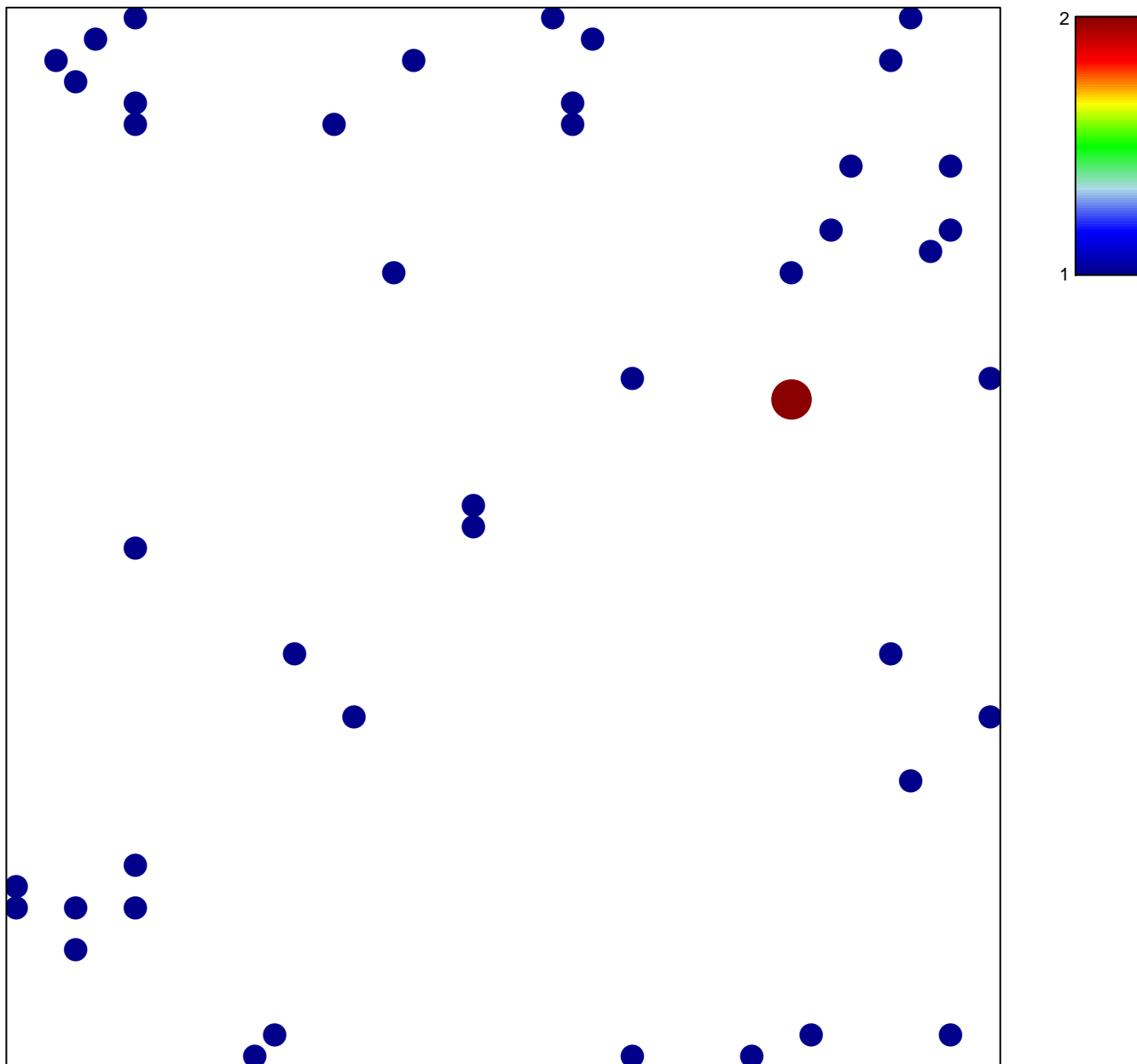
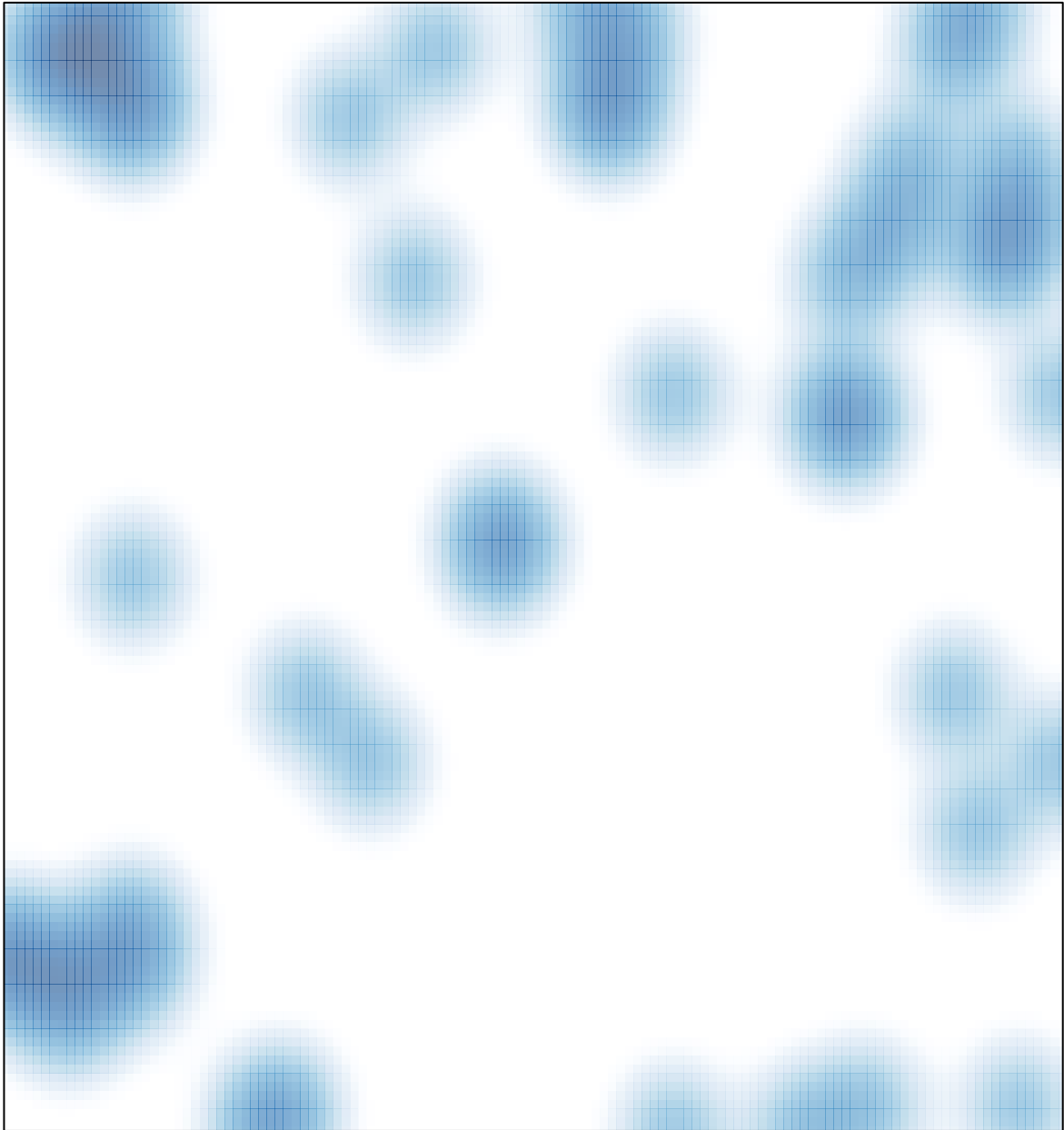


# manganese ion binding



# features = 45  
chi-square p = 0.81

# manganese ion binding



# features = 45 , max = 2