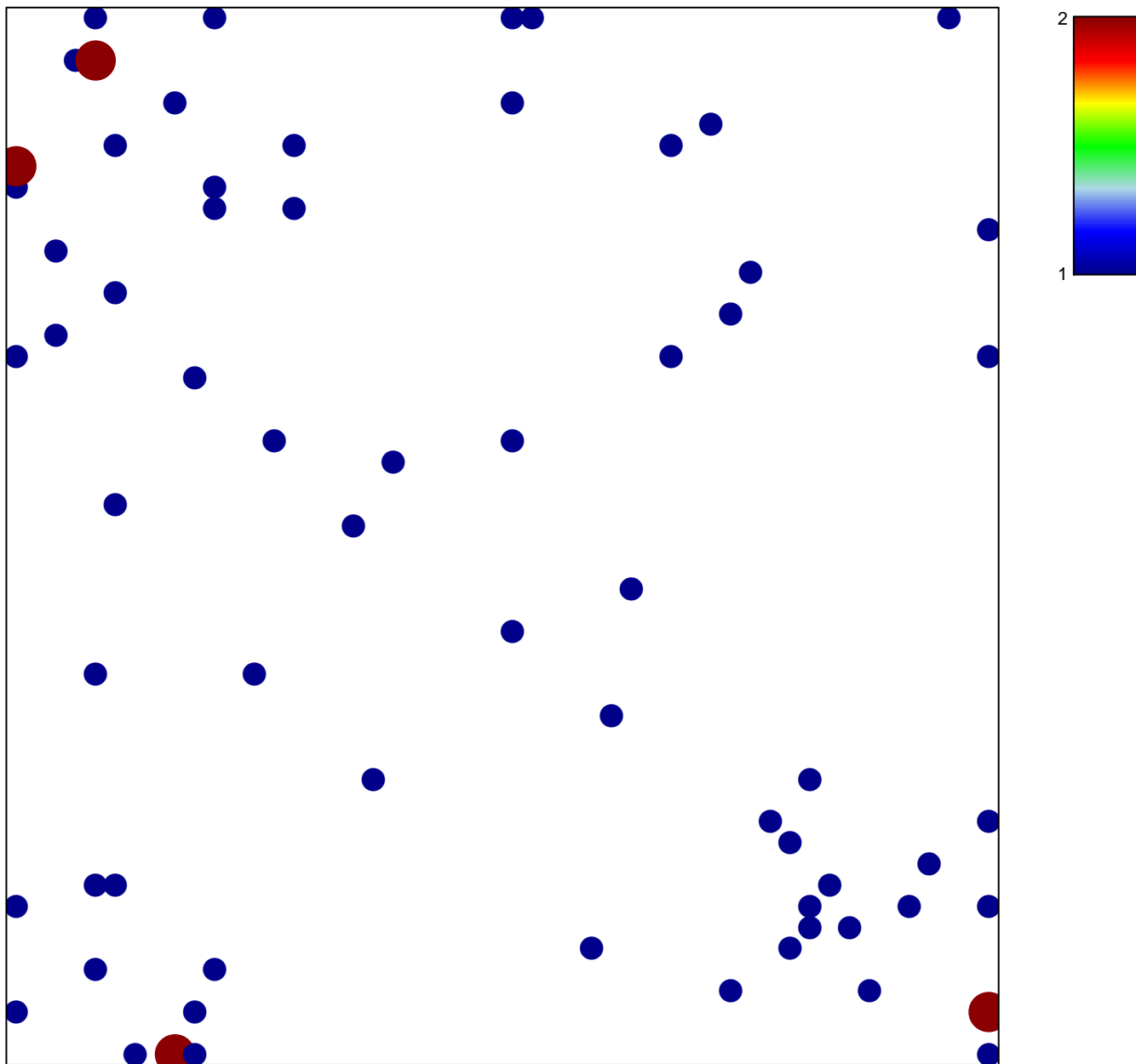
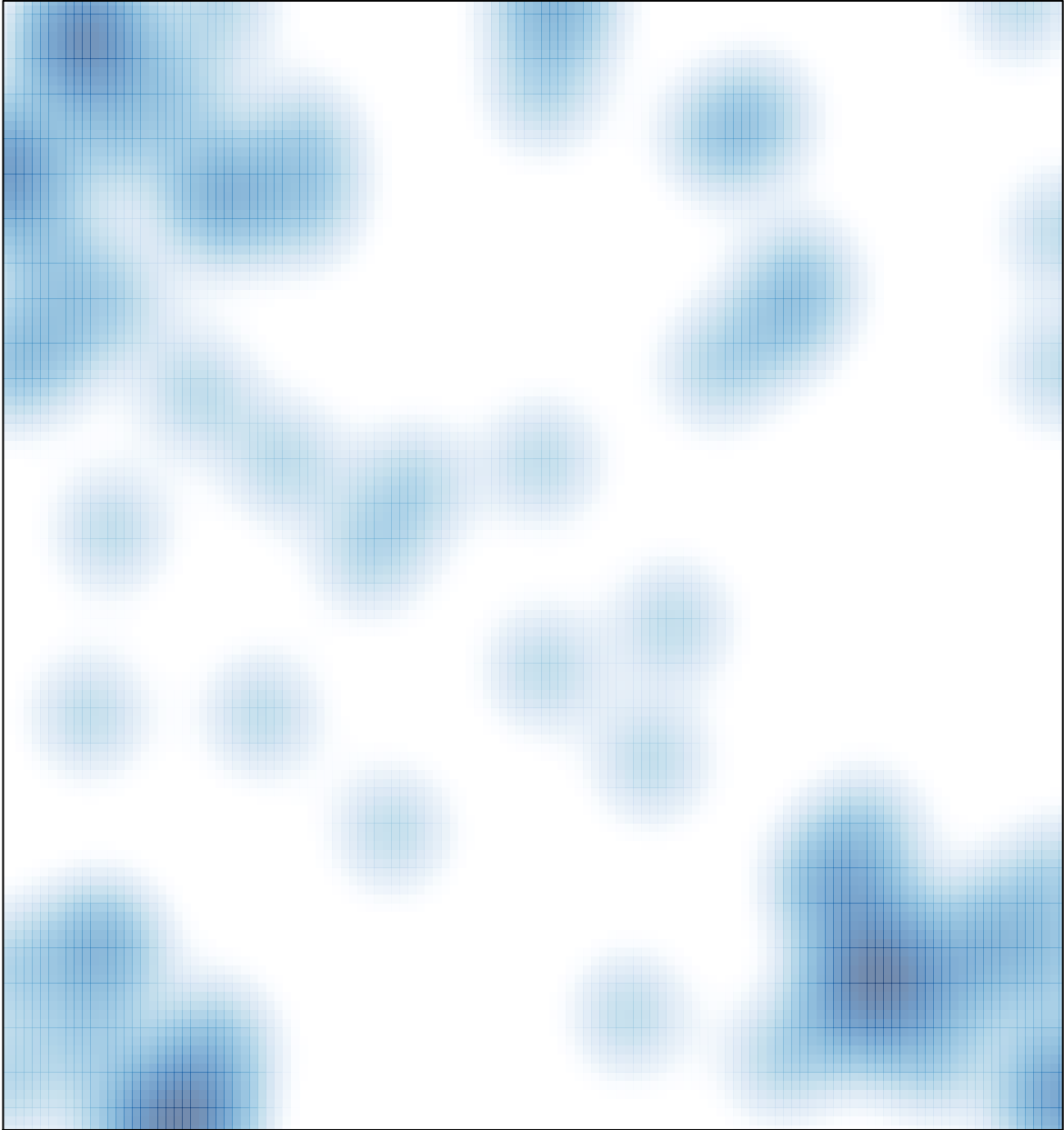


# phospholipid binding



# features = 70  
chi-square p = 0.8

# phospholipid binding



# features = 70 , max = 2