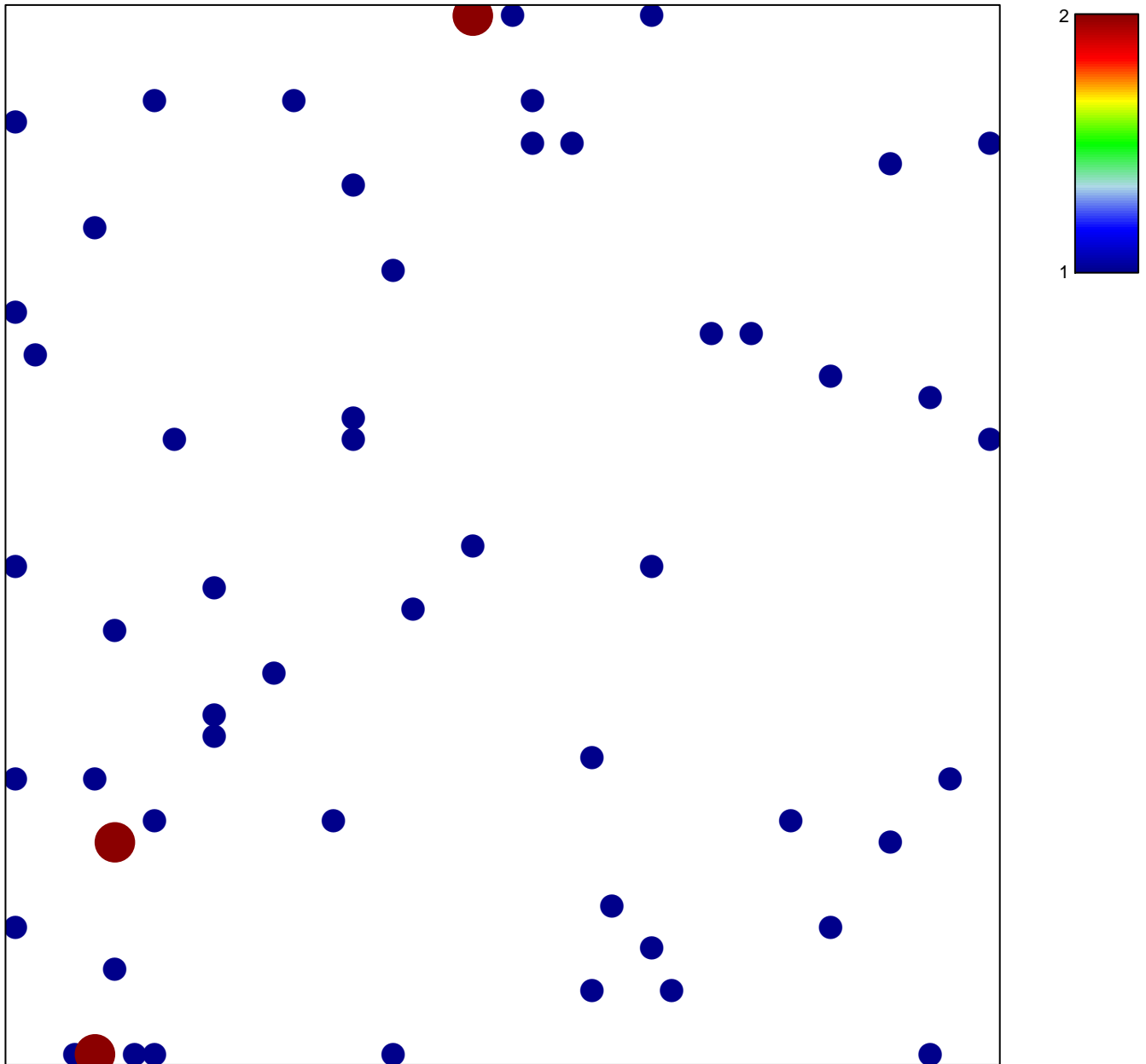
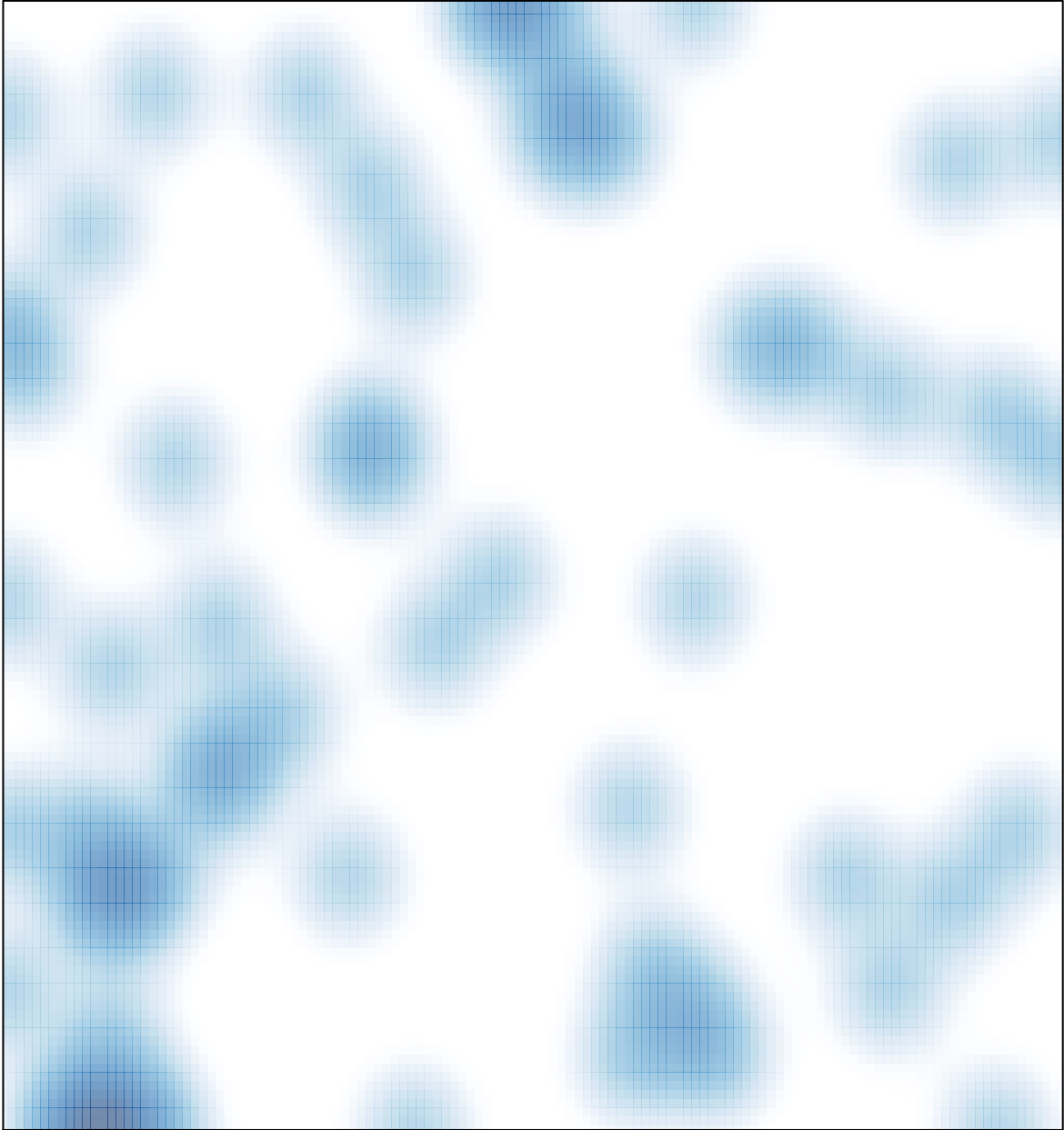


# protein heterooligomerization



# features = 58  
chi-square p = 0.81

# protein heterooligomerization



# features = 58 , max = 2