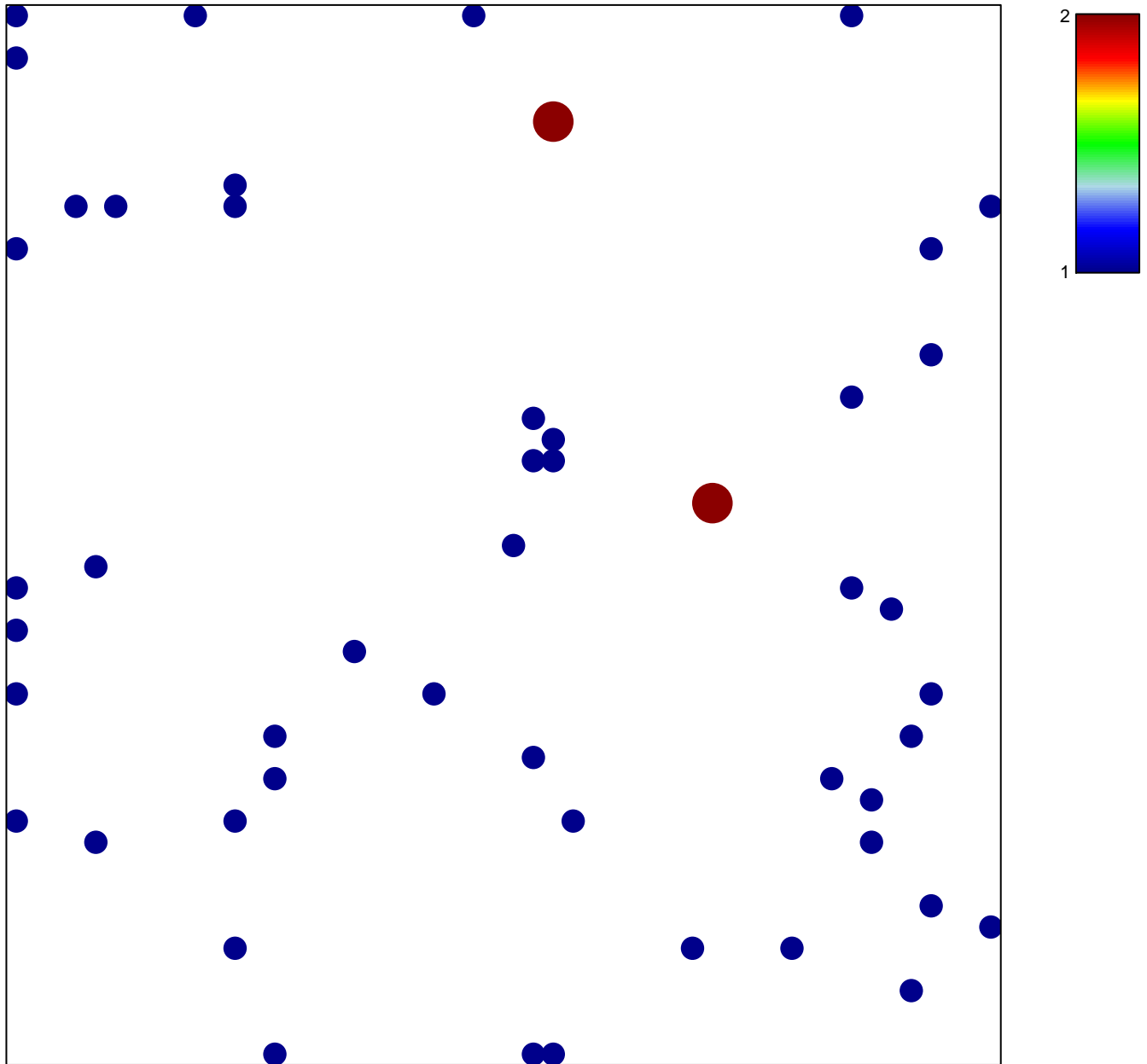
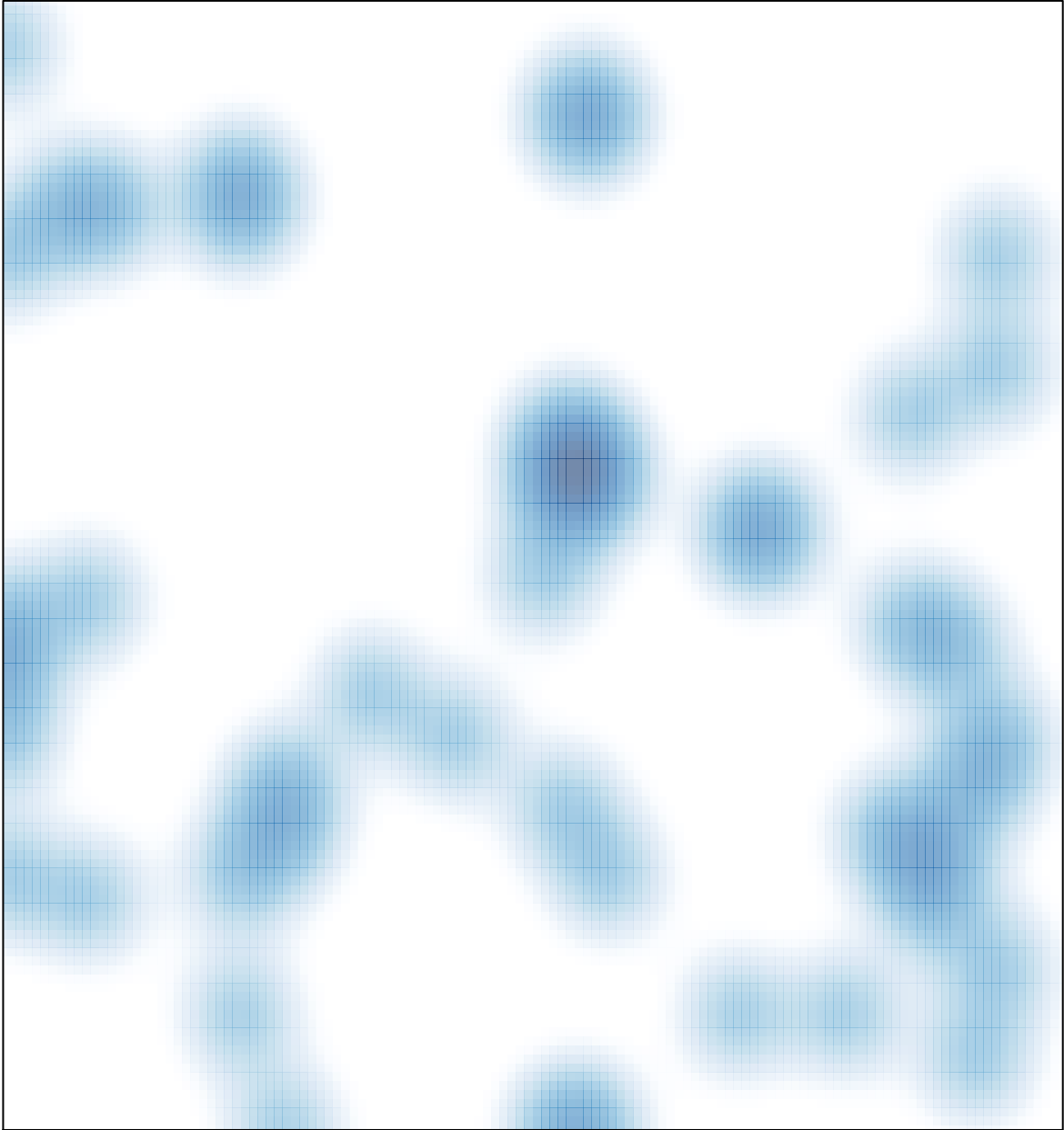


# protein heterooligomerization



# features = 52  
chi-square p = 0.79

# protein heterooligomerization



# features = 52 , max = 2