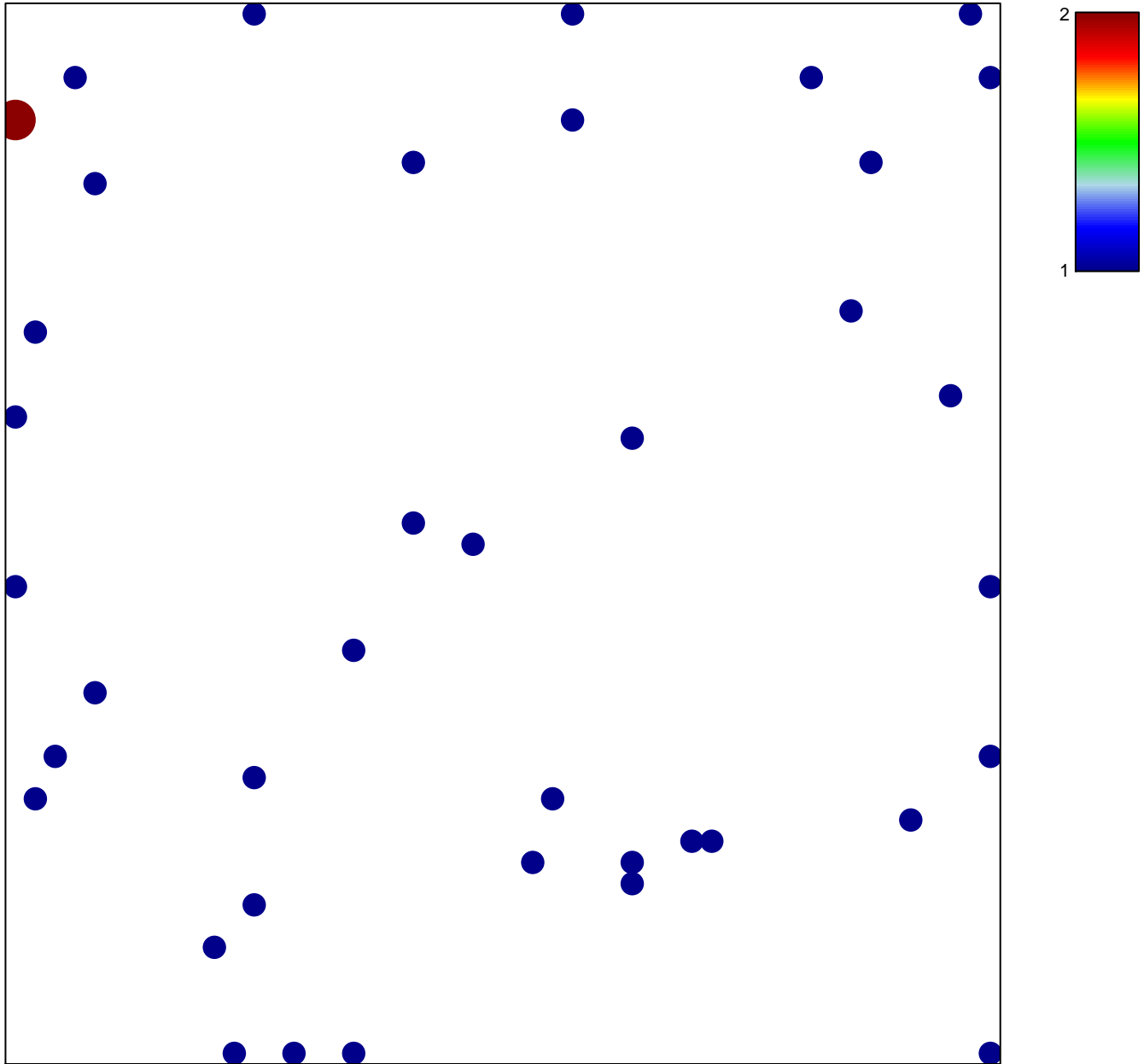
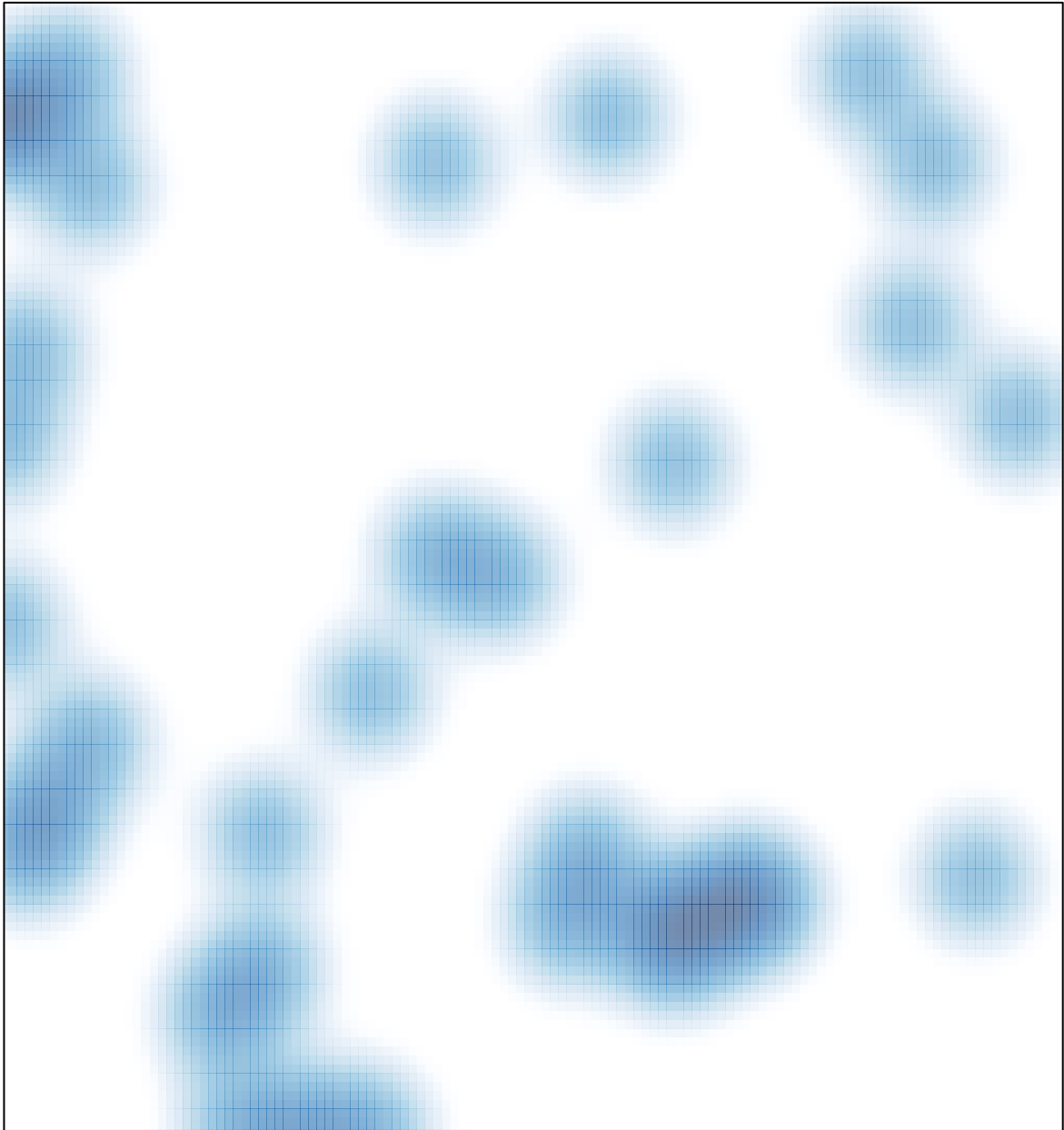


# heat shock protein binding



# features = 40  
chi-square p = 0.82

# heat shock protein binding



# features = 40 , max = 2