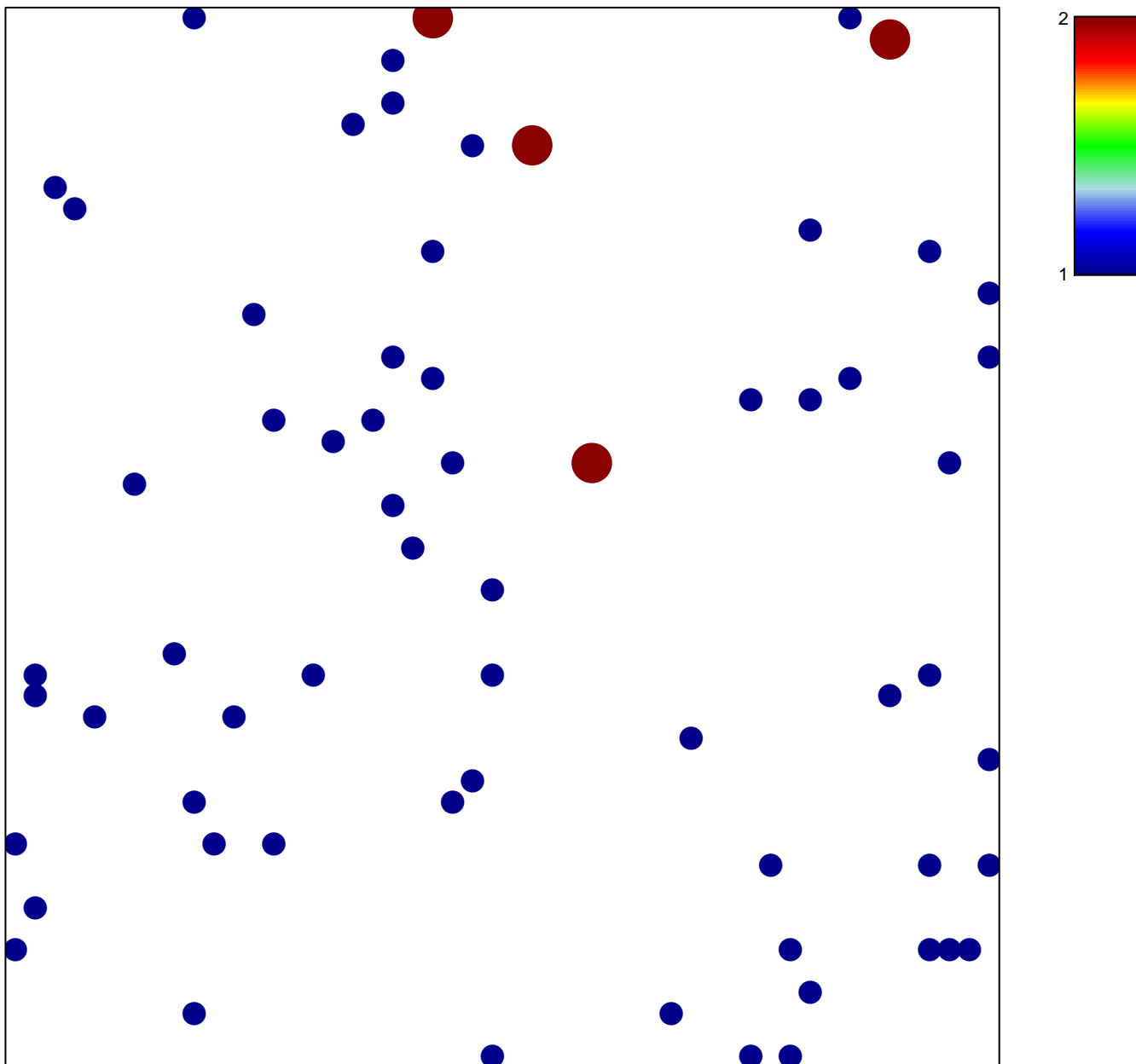
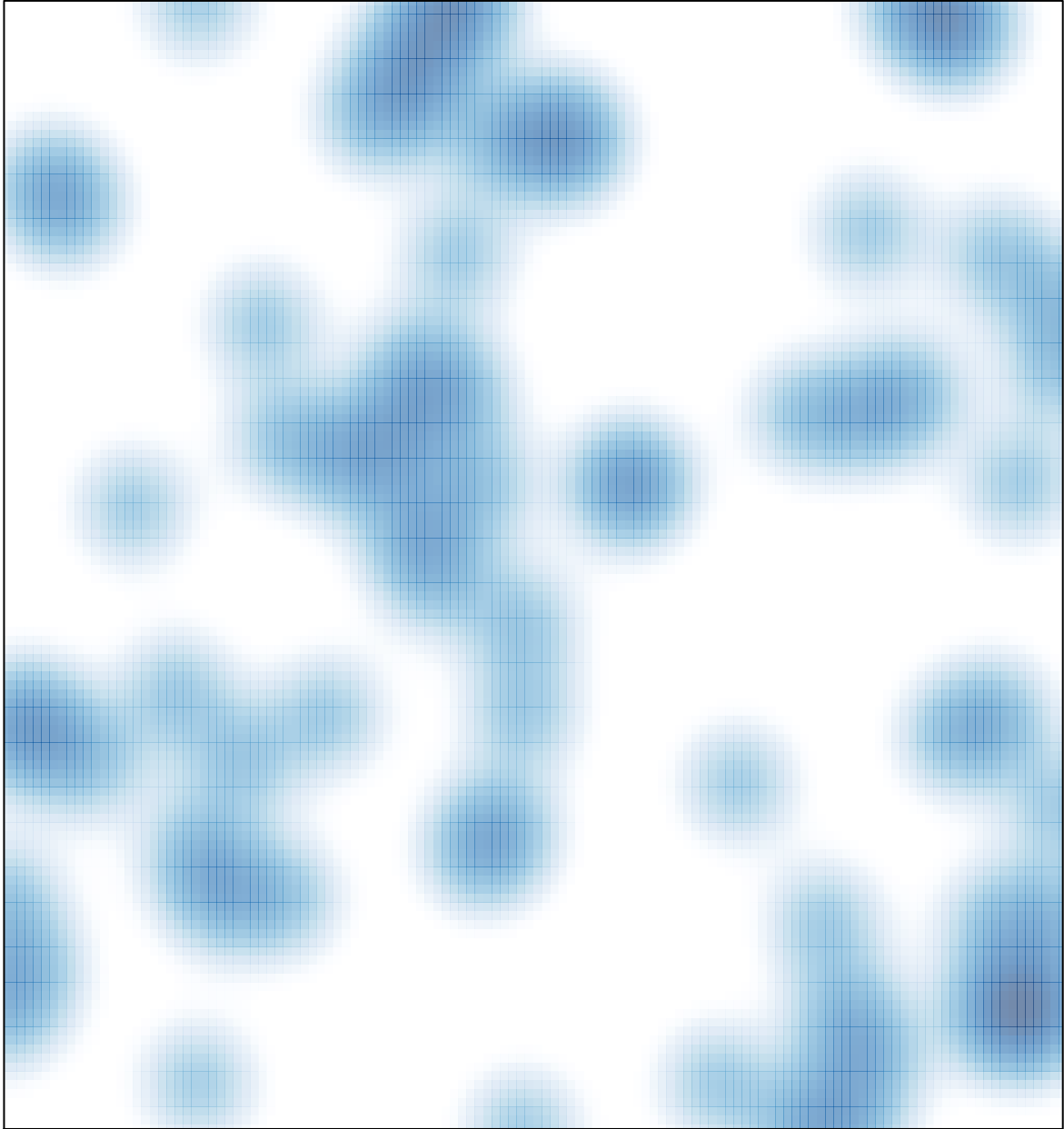


# histone deacetylase binding



# features = 68  
chi-square p = 0.83

# histone deacetylase binding



# features = 68 , max = 2