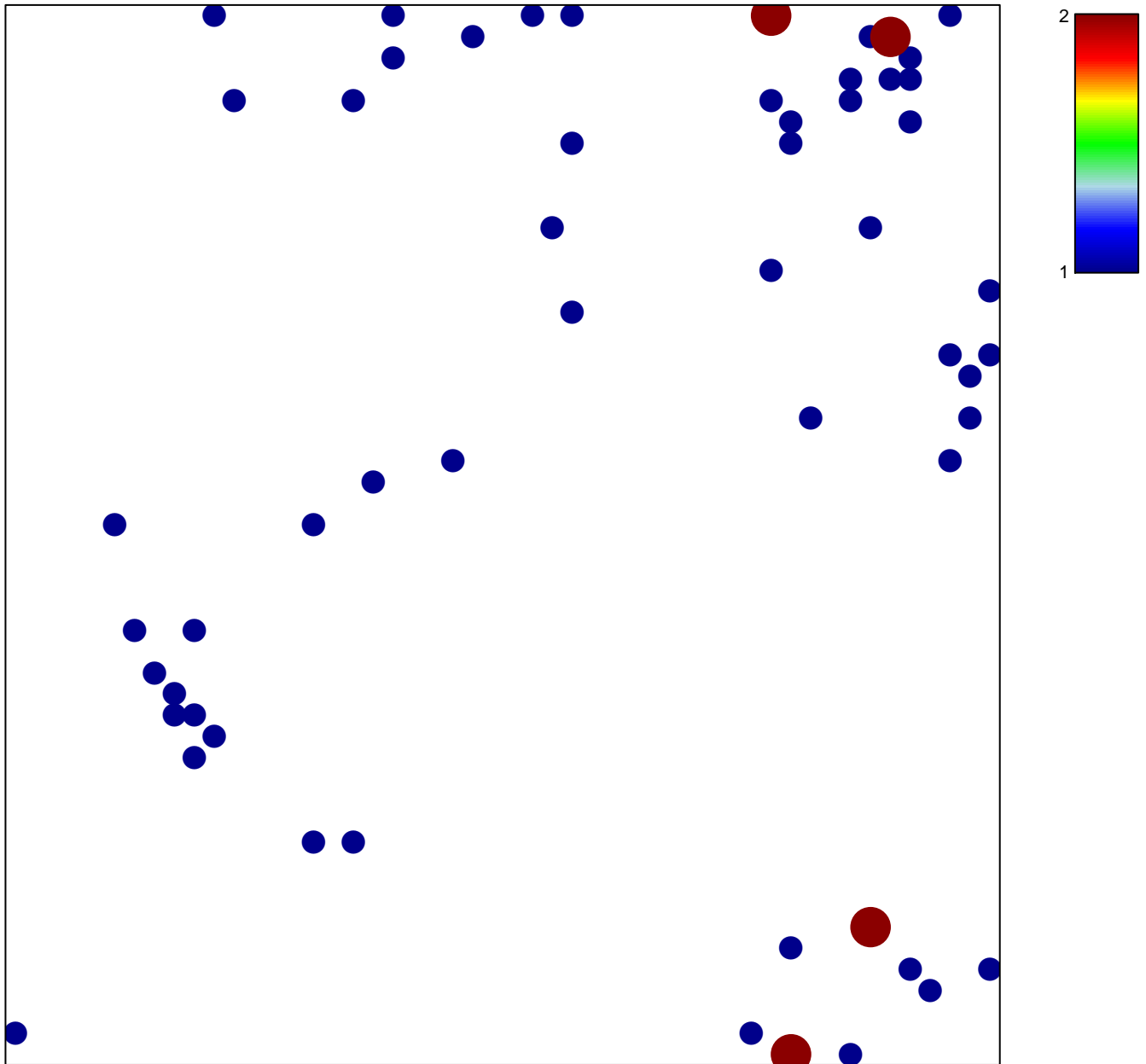
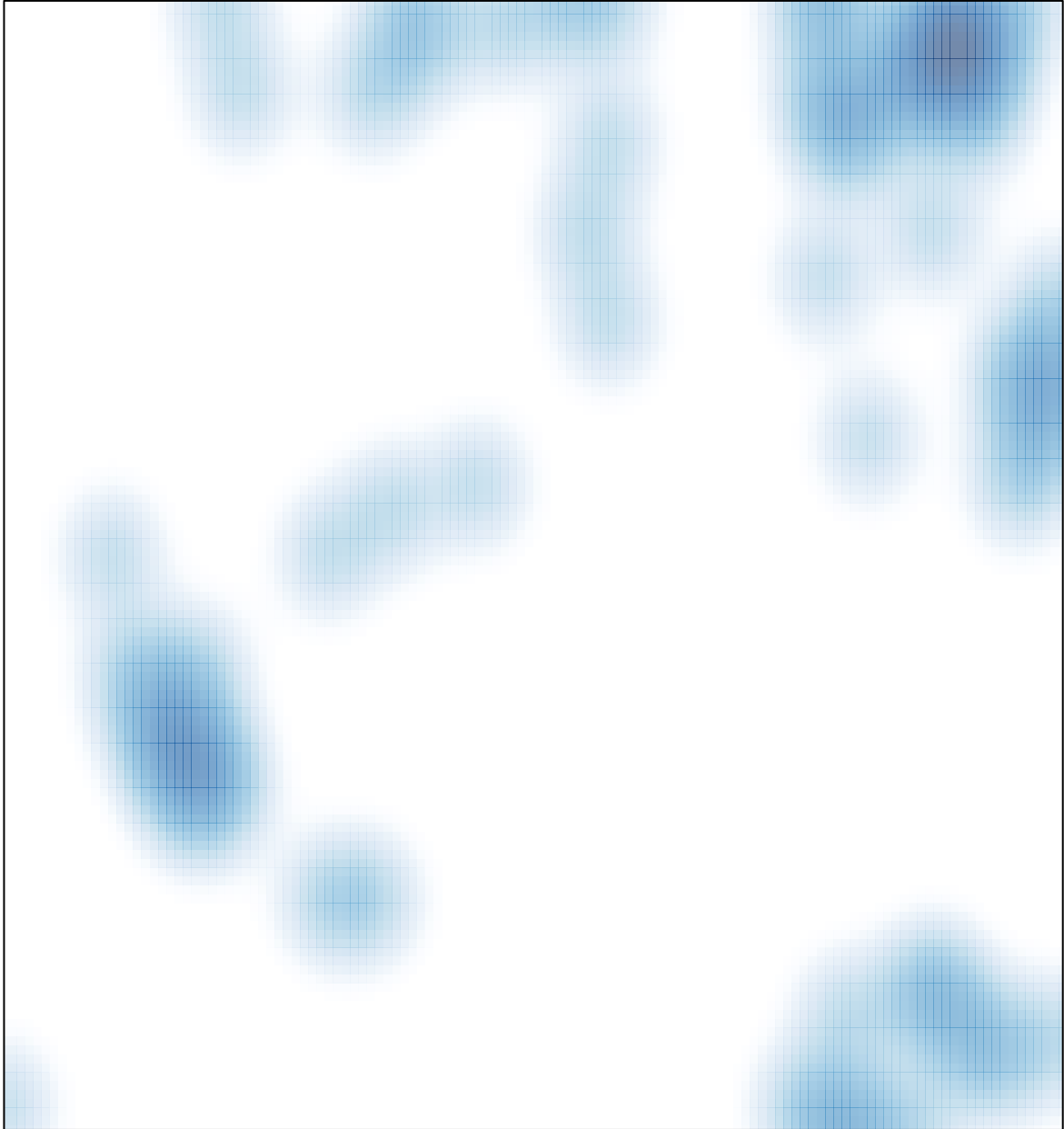


histone binding



features = 60
chi-square p = 0.8

histone binding



features = 60 , max = 2