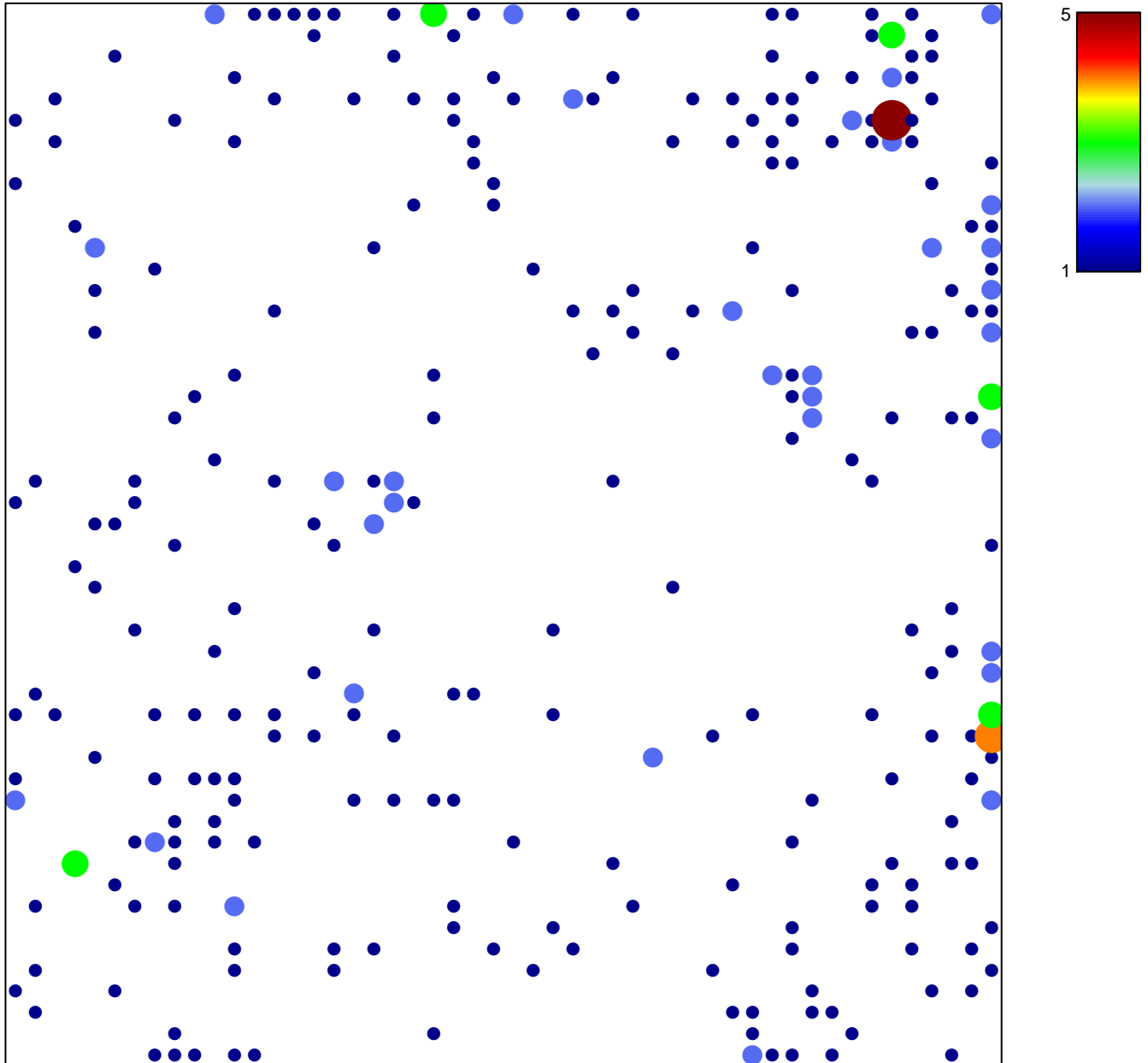
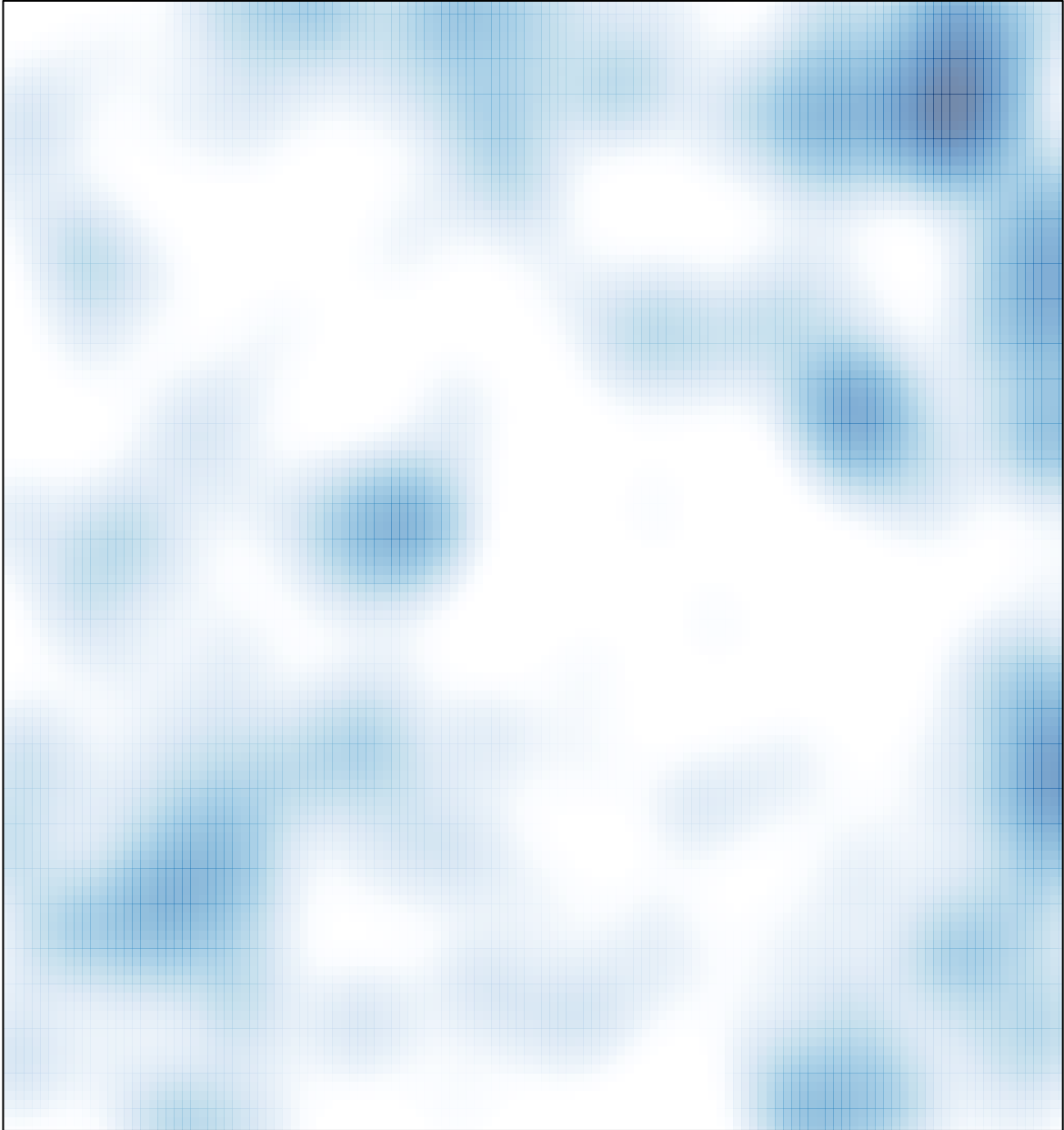


chromatin binding



features = 318
chi-square p = 0.72

chromatin binding



features = 318 , max = 5