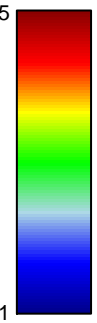
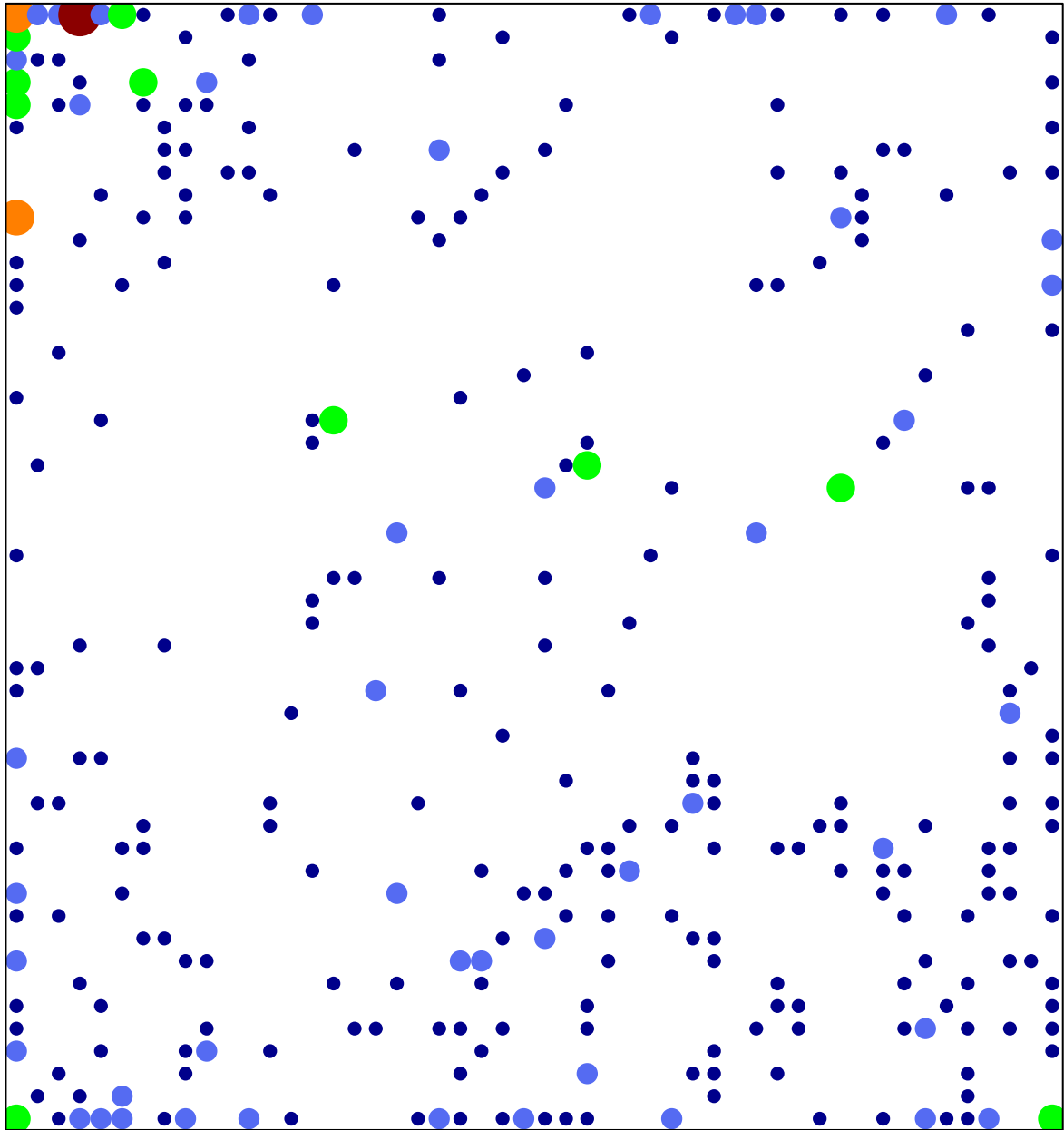
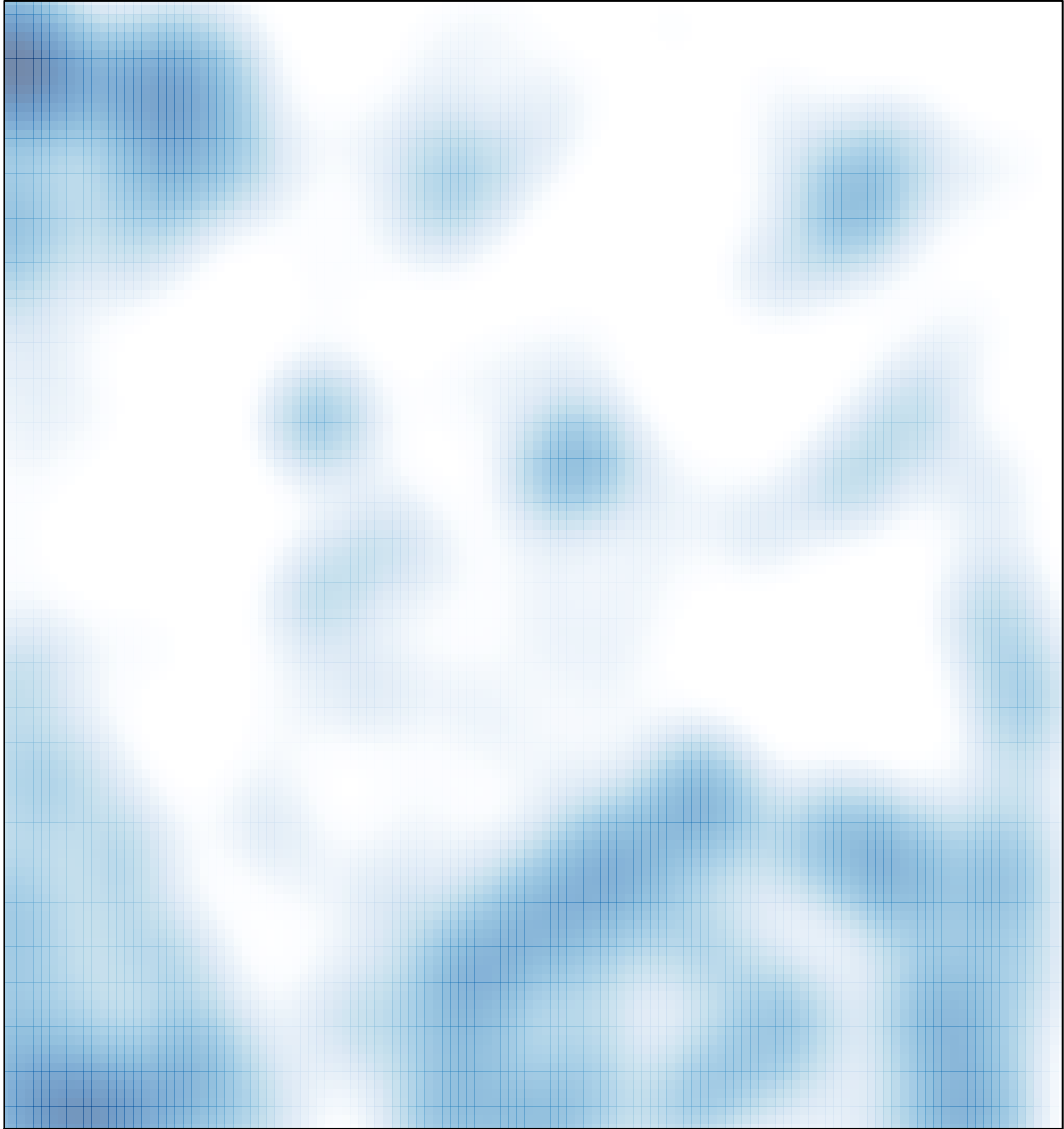


chromatin binding



features = 376
chi-square p = 0.62

chromatin binding



features = 376 , max = 5