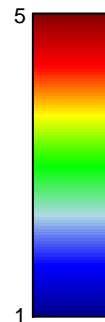
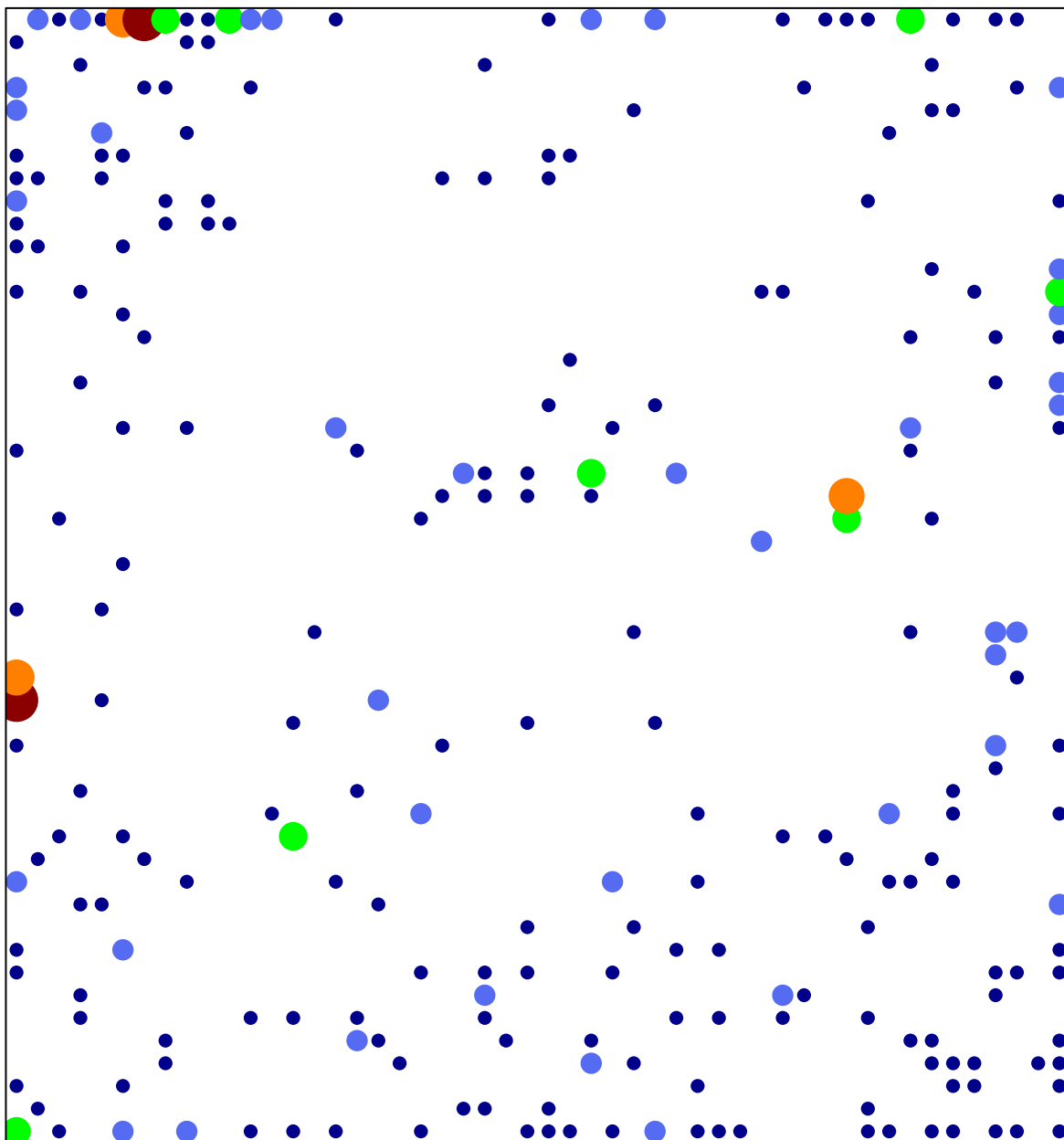
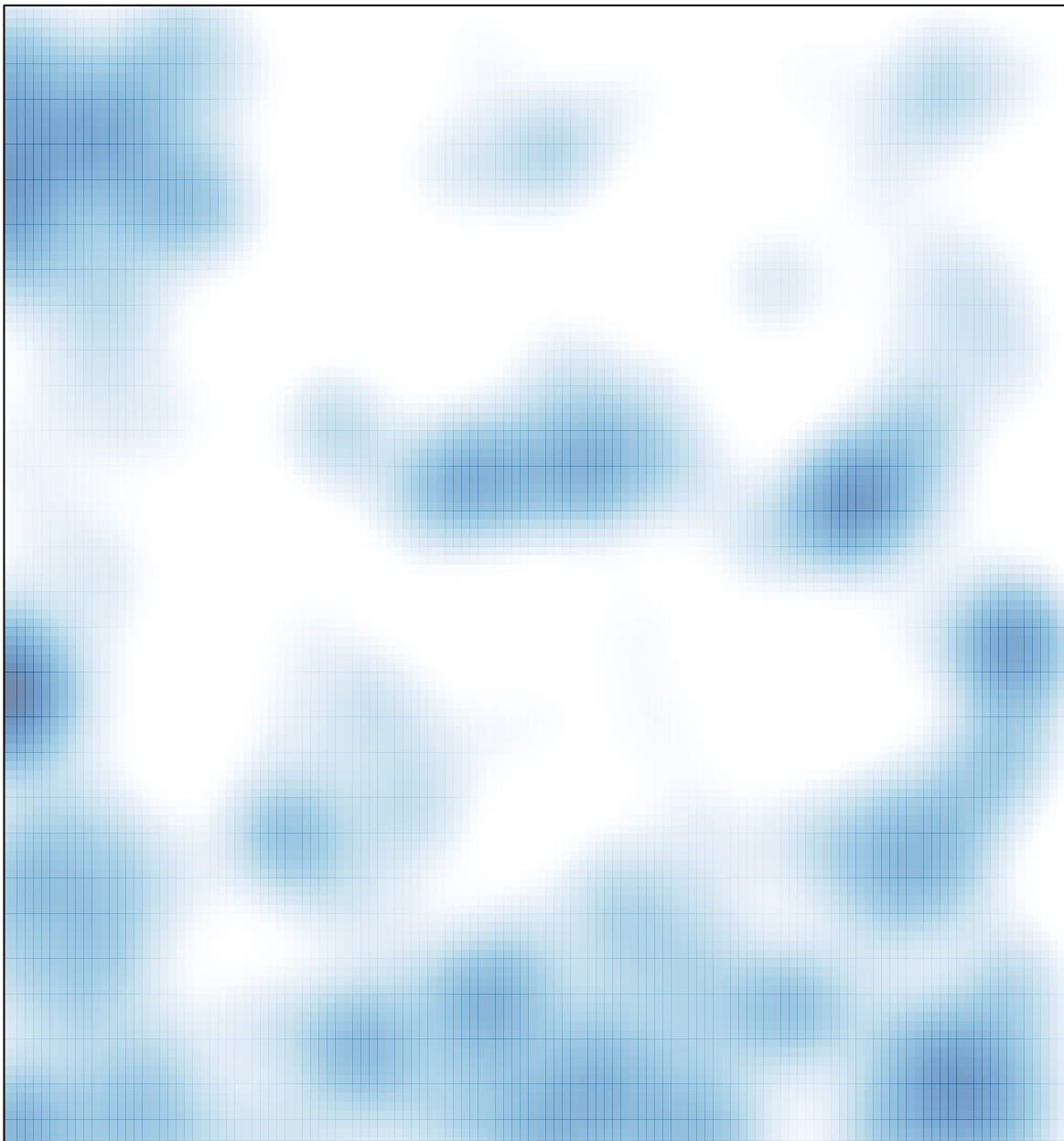


protein kinase binding



features = 316
chi-square p = 0.63

protein kinase binding



features = 316 , max = 5