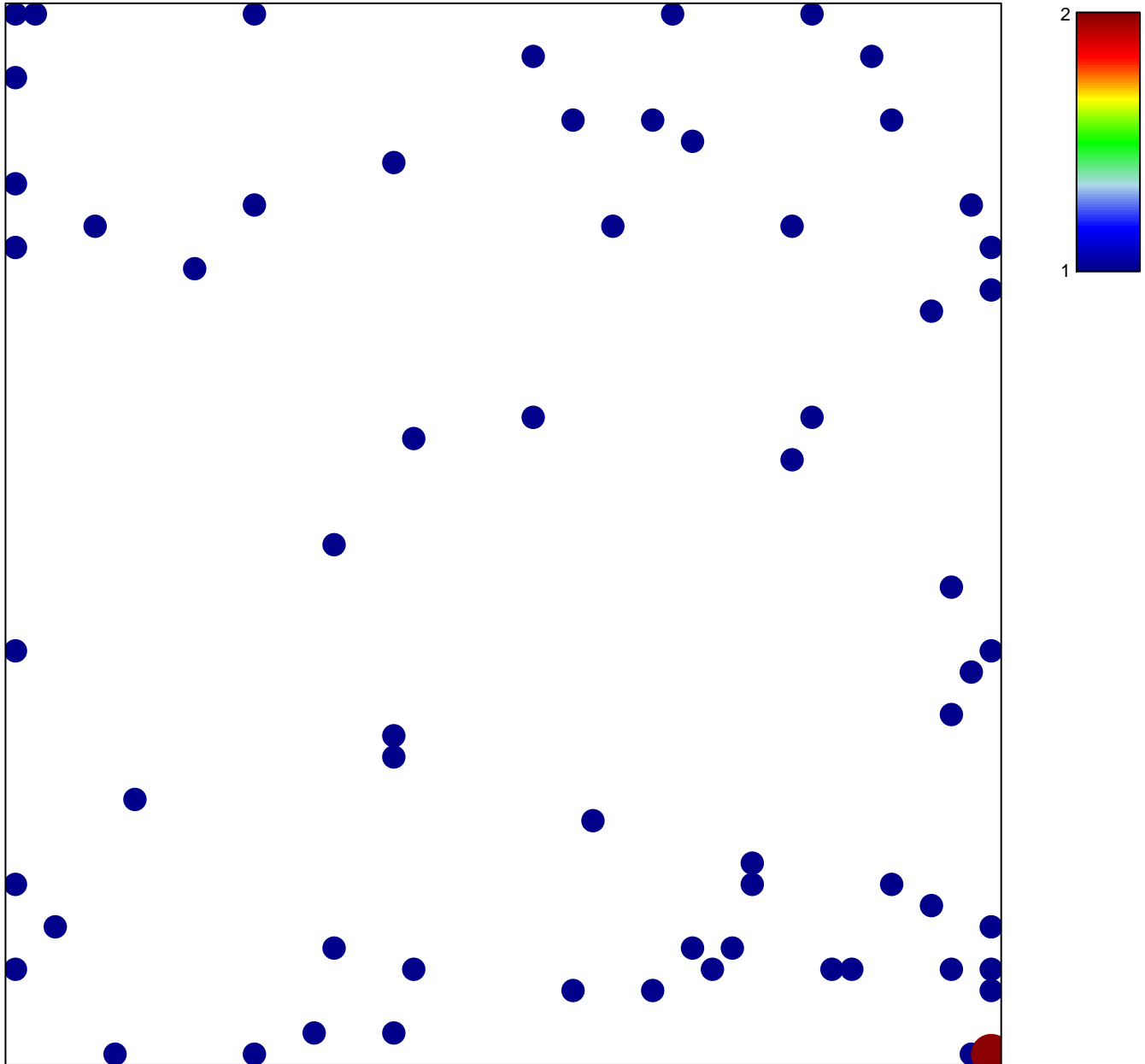
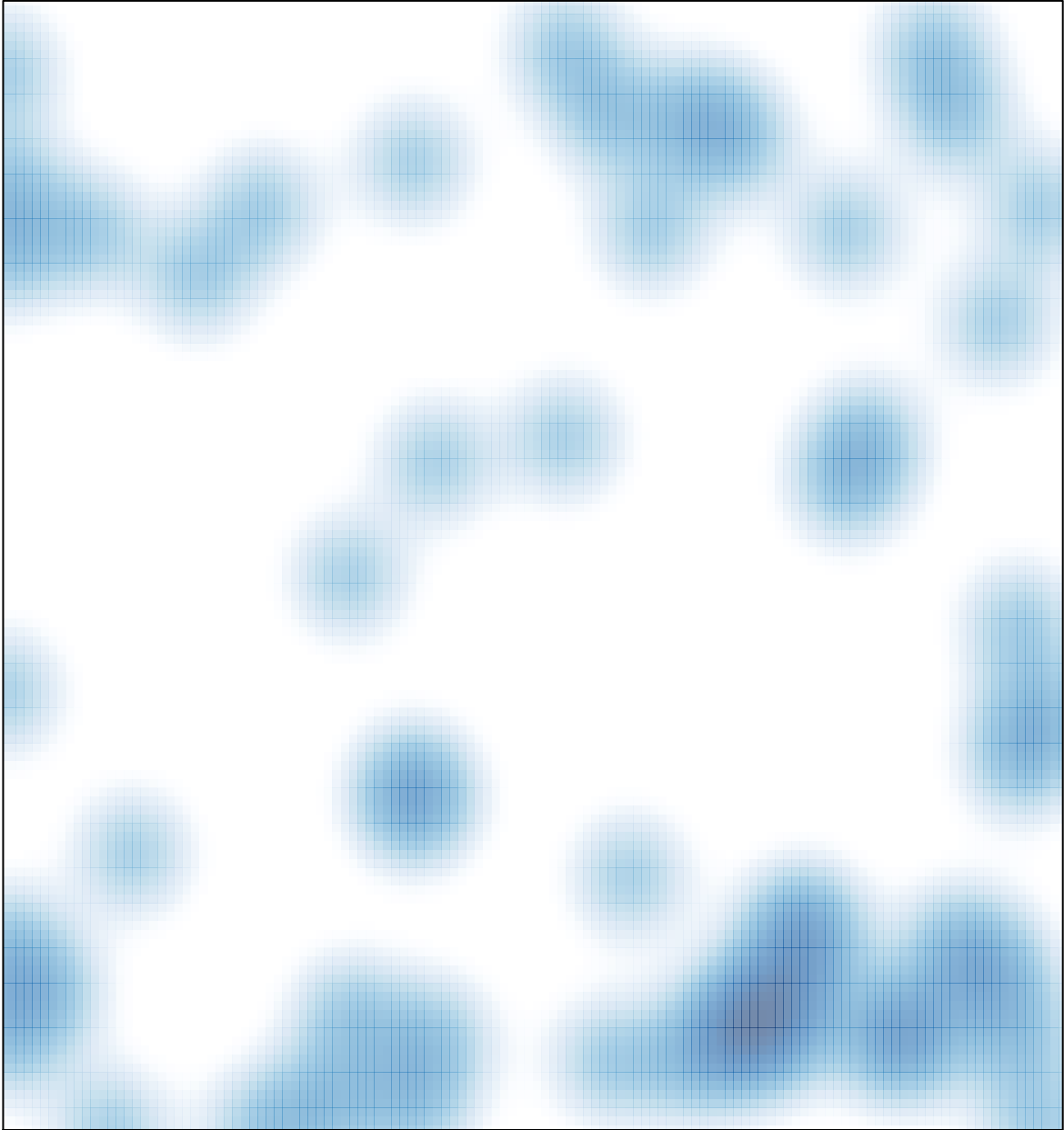


protease binding



features = 65
chi-square p = 0.79

protease binding



features = 65 , max = 2