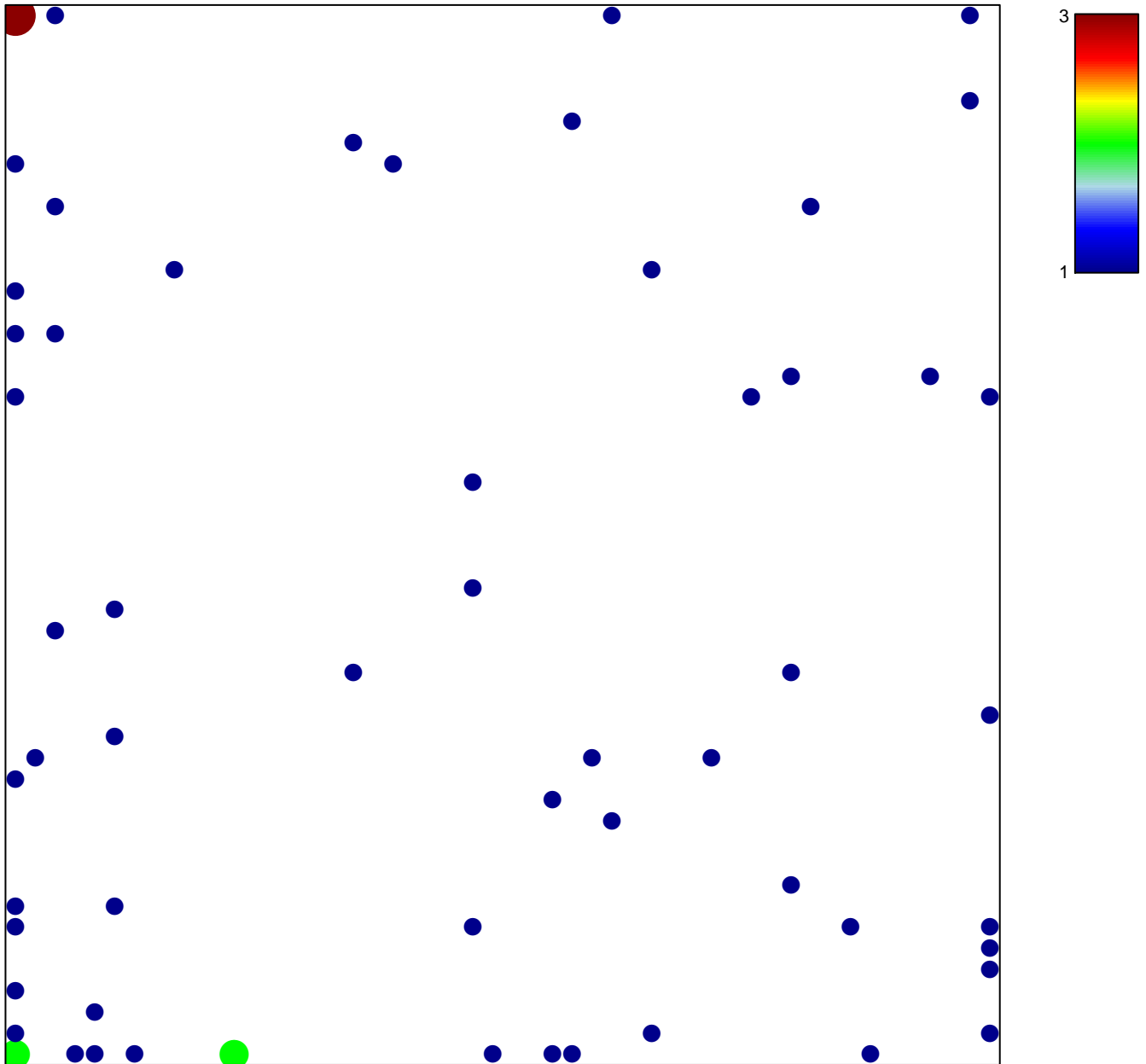
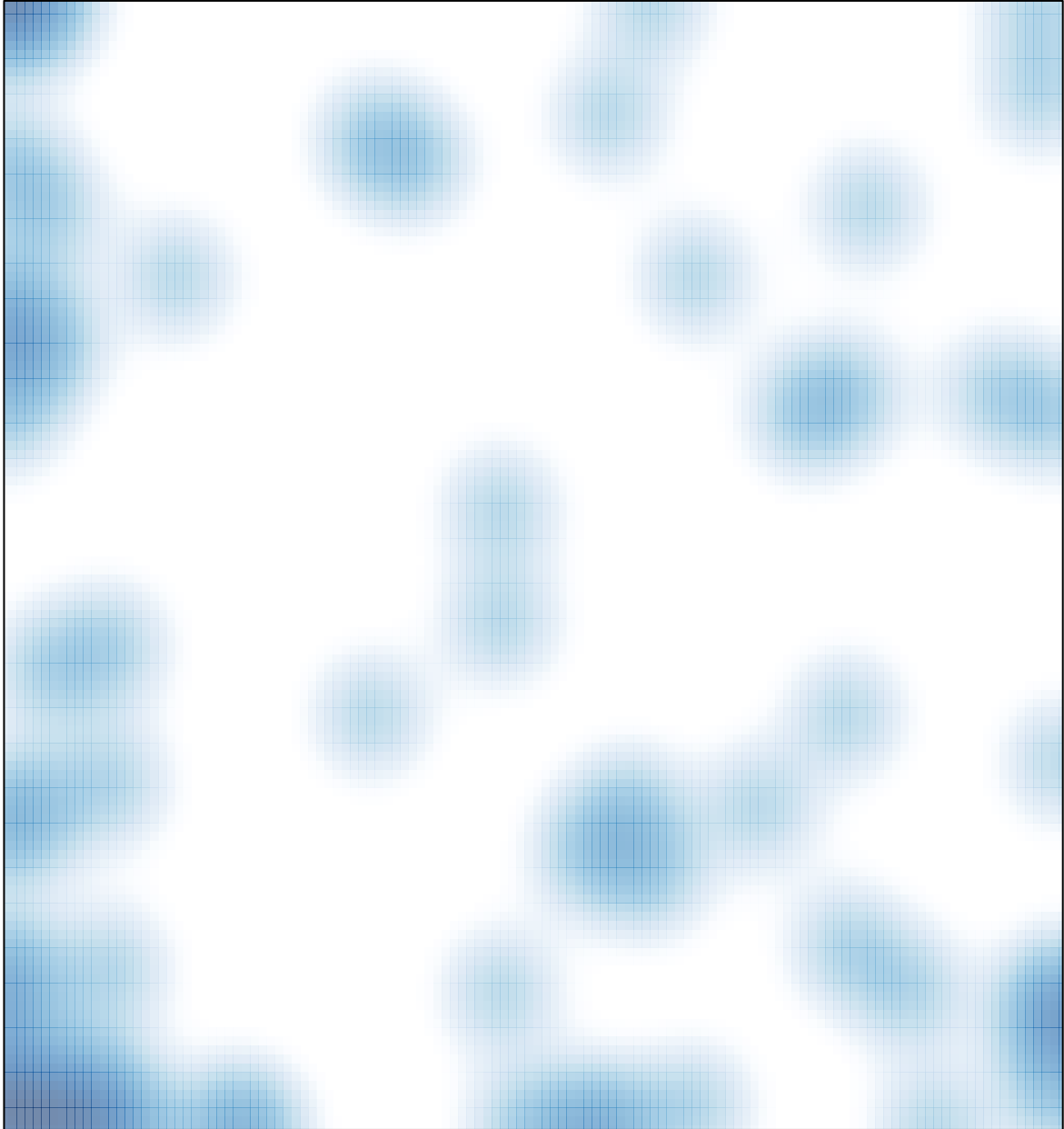


protease binding



features = 62
chi-square p = 0.72

protease binding



features = 62 , max = 3