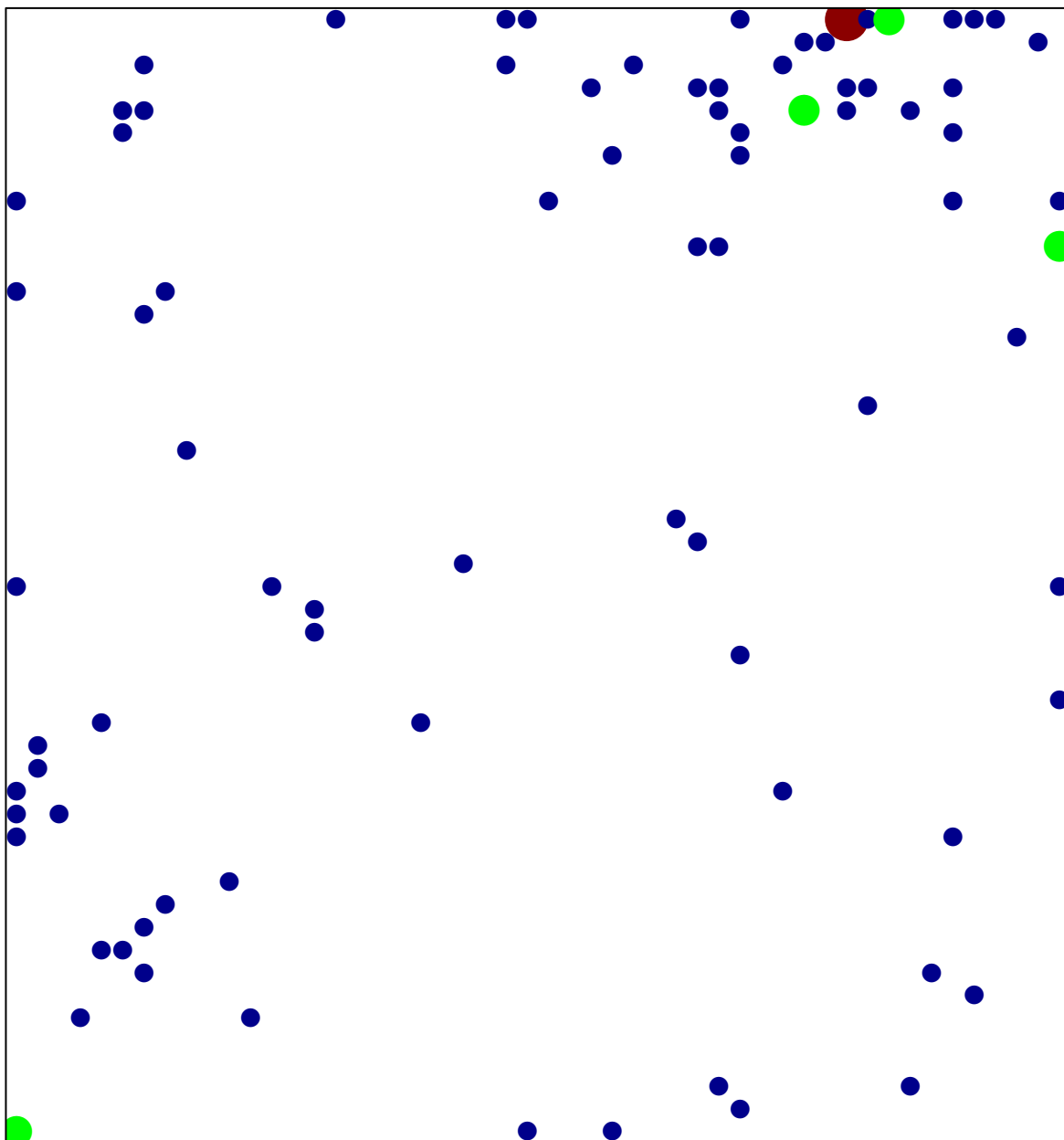
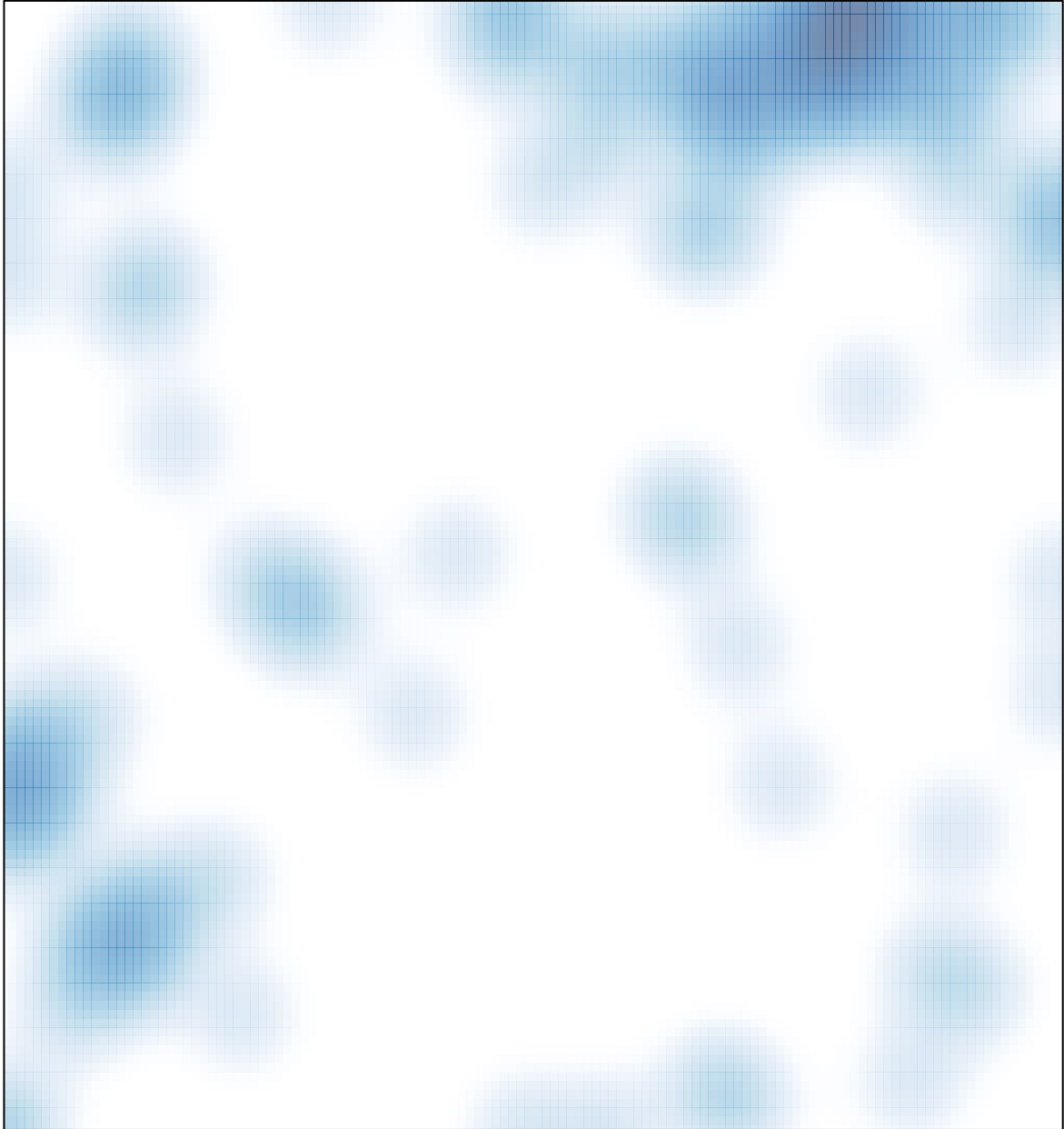


unfolded protein binding



features = 89
chi-square p = 0.77

unfolded protein binding



features = 89 , max = 3