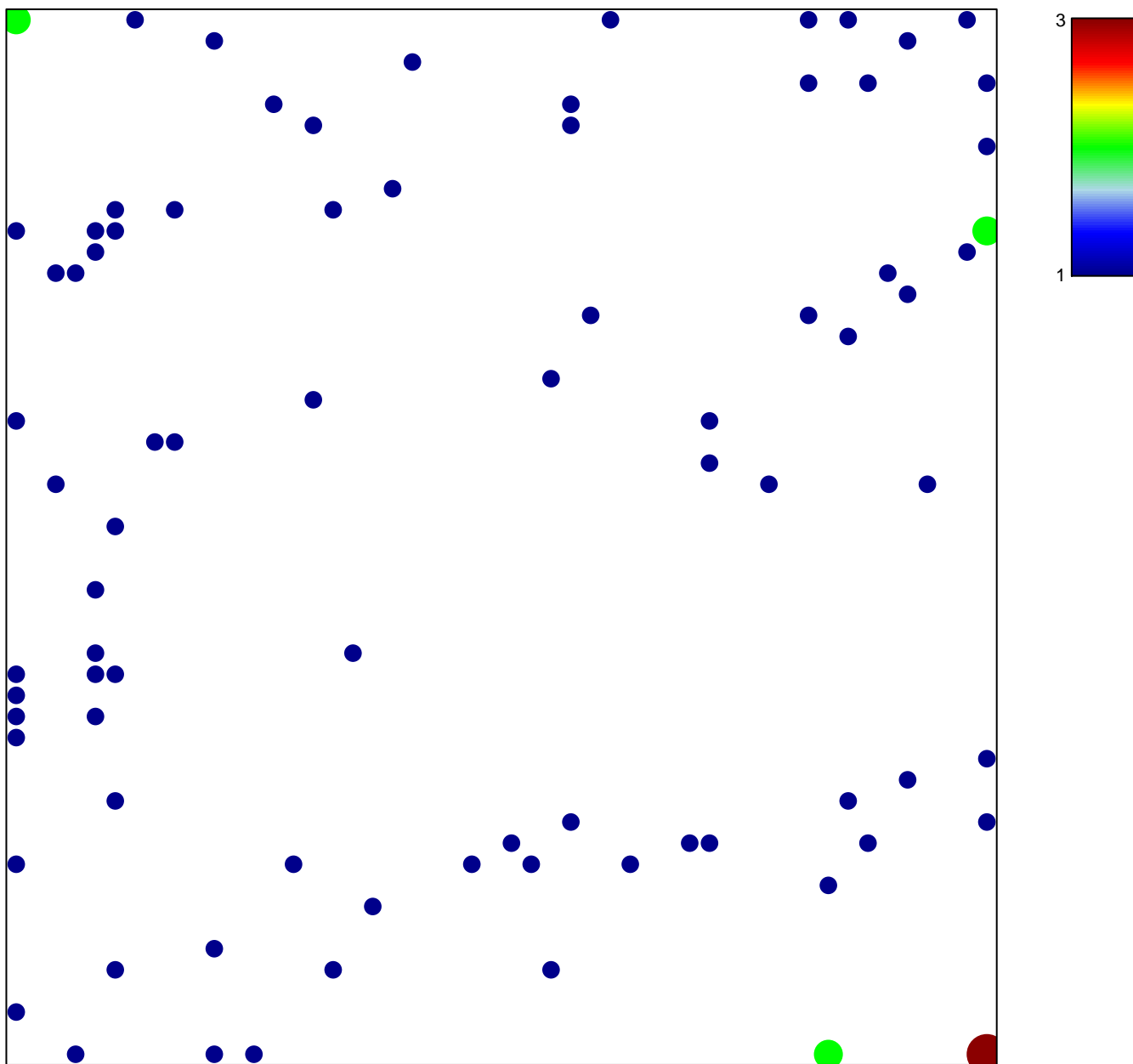
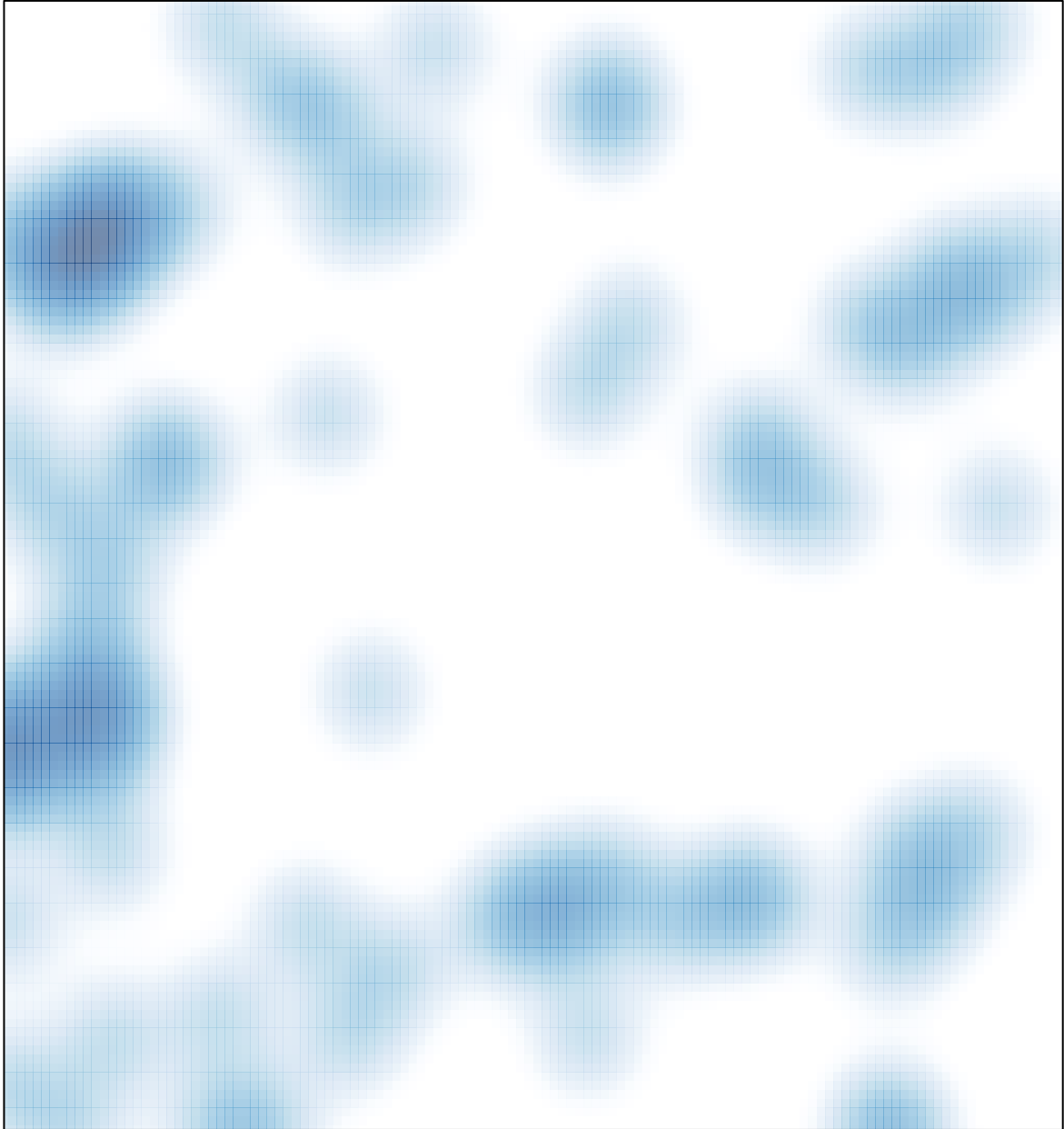


unfolded protein binding



features = 87
chi-square p = 0.77

unfolded protein binding



features = 87 , max = 3