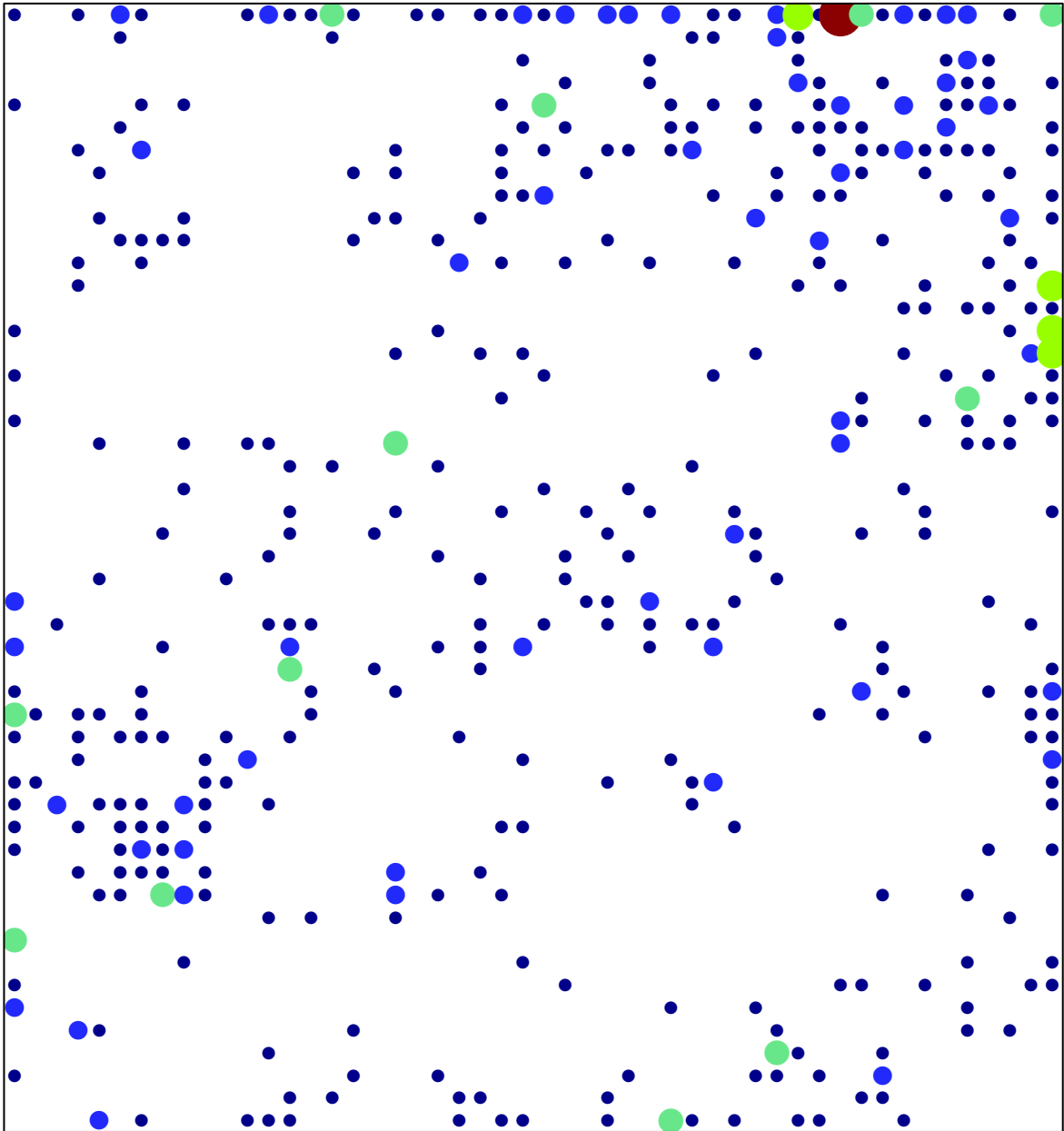
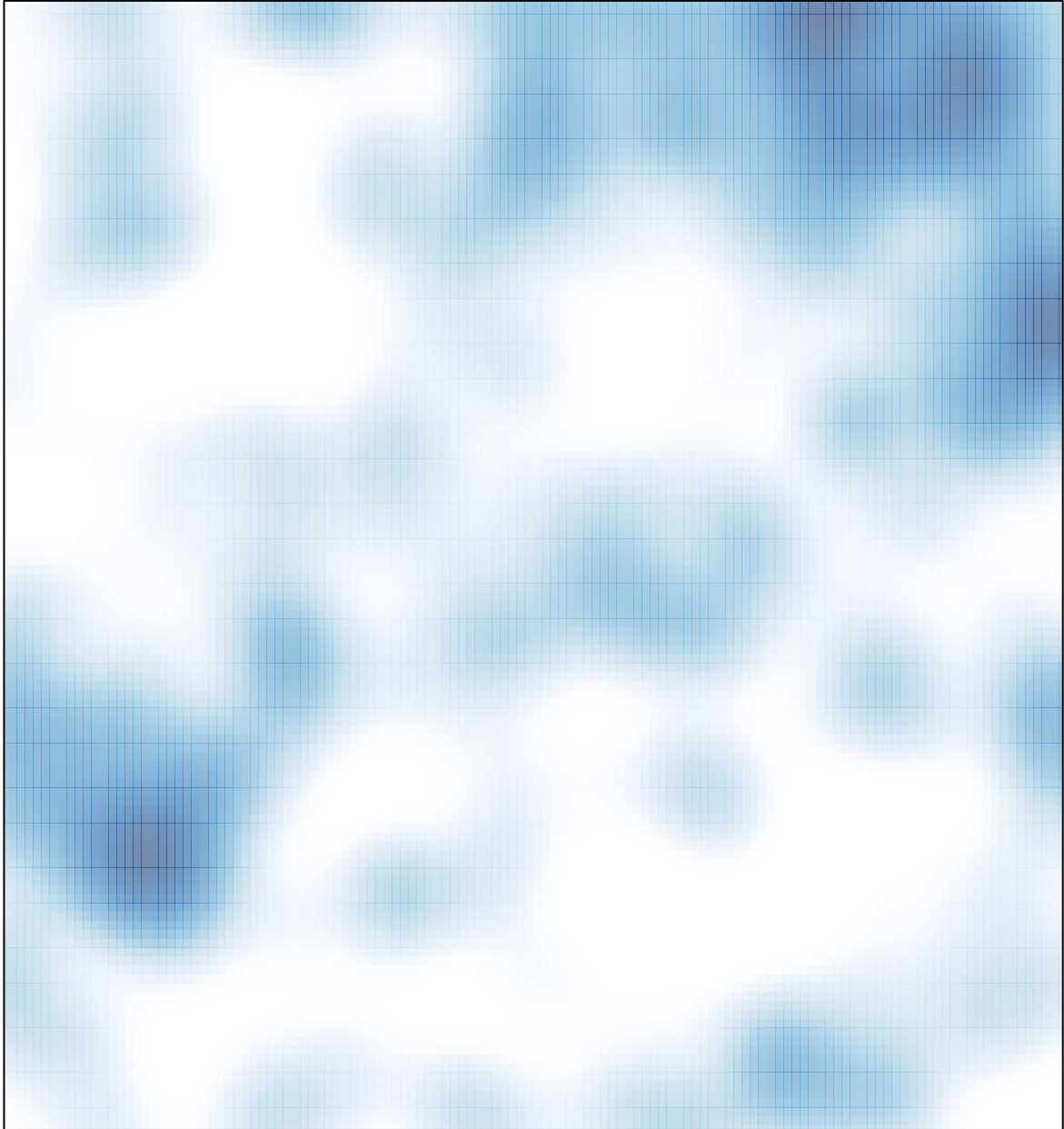


nucleotide binding



features = 506
chi-square p = 0.66

nucleotide binding



features = 506 , max = 6