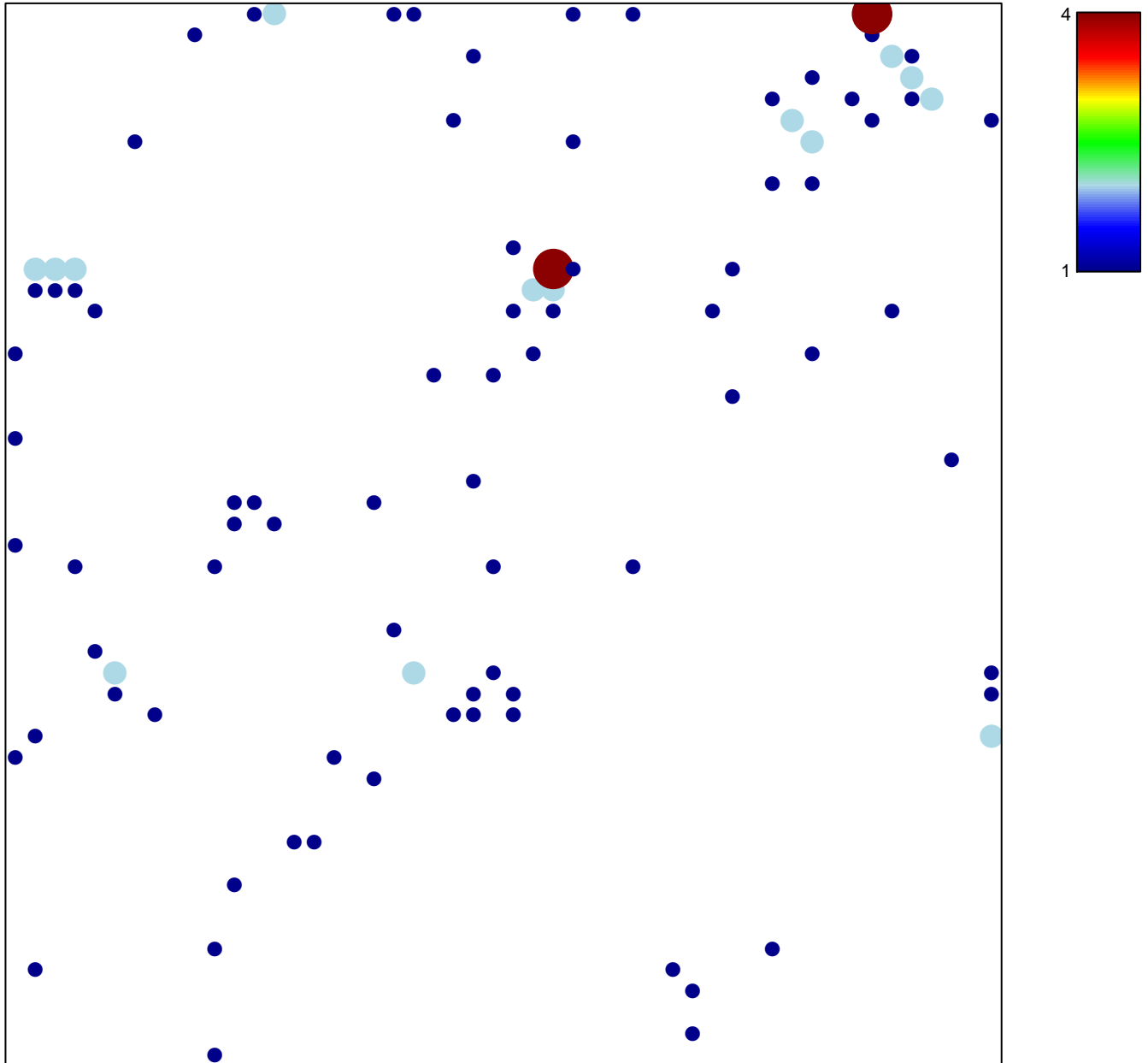
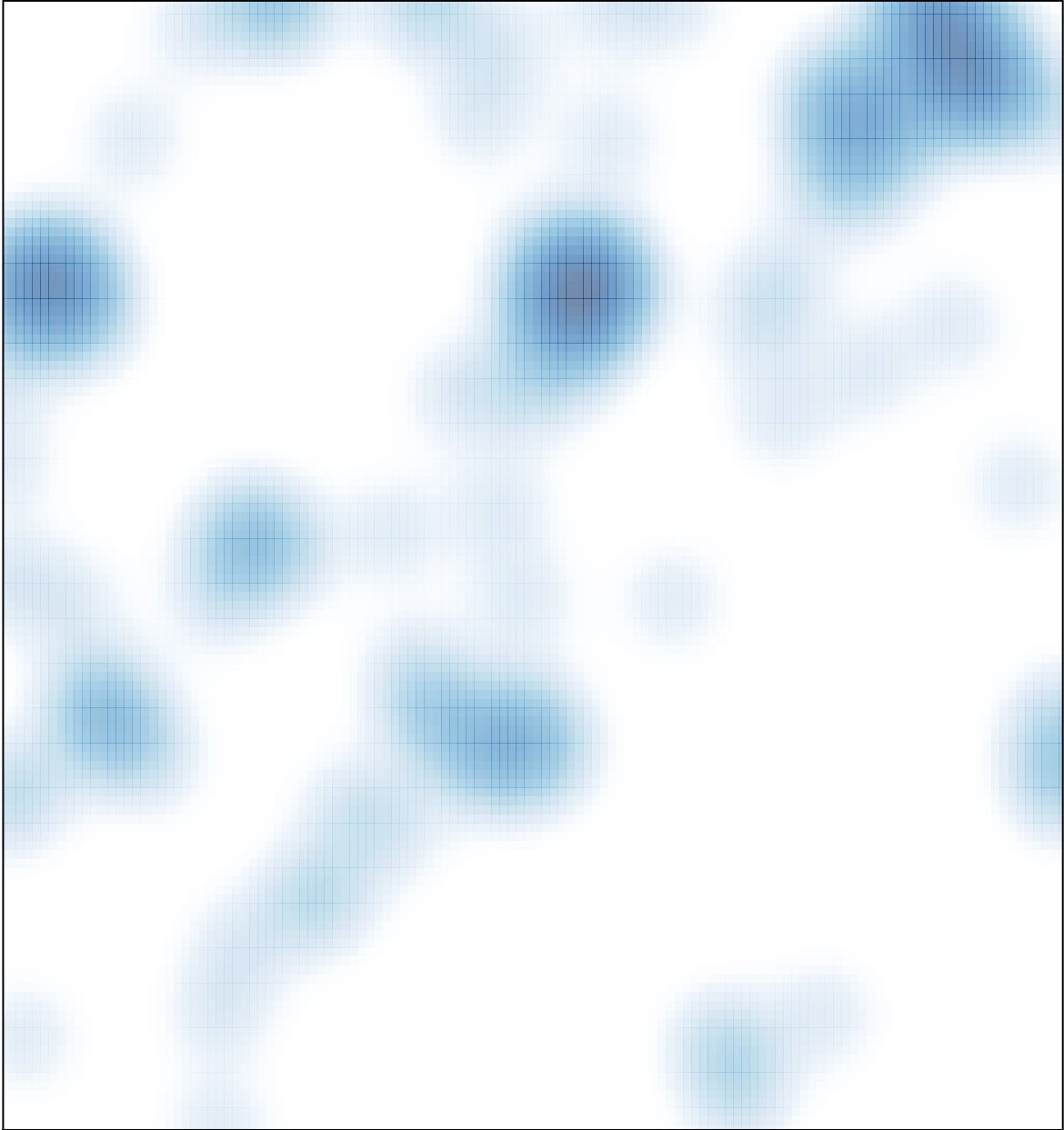


nucleosome assembly



features = 112
chi-square p = 0.77

nucleosome assembly



features = 112 , max = 4