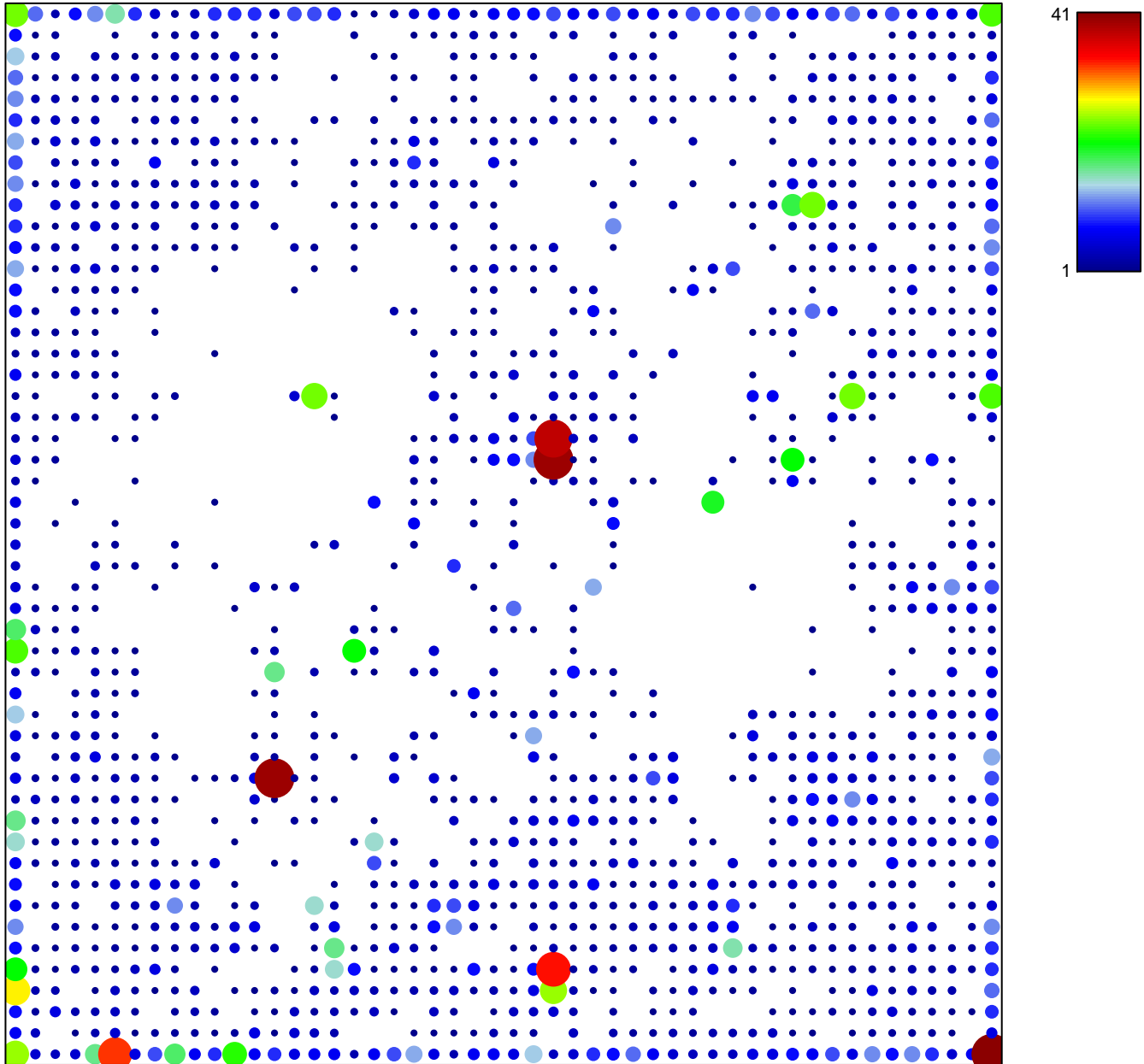
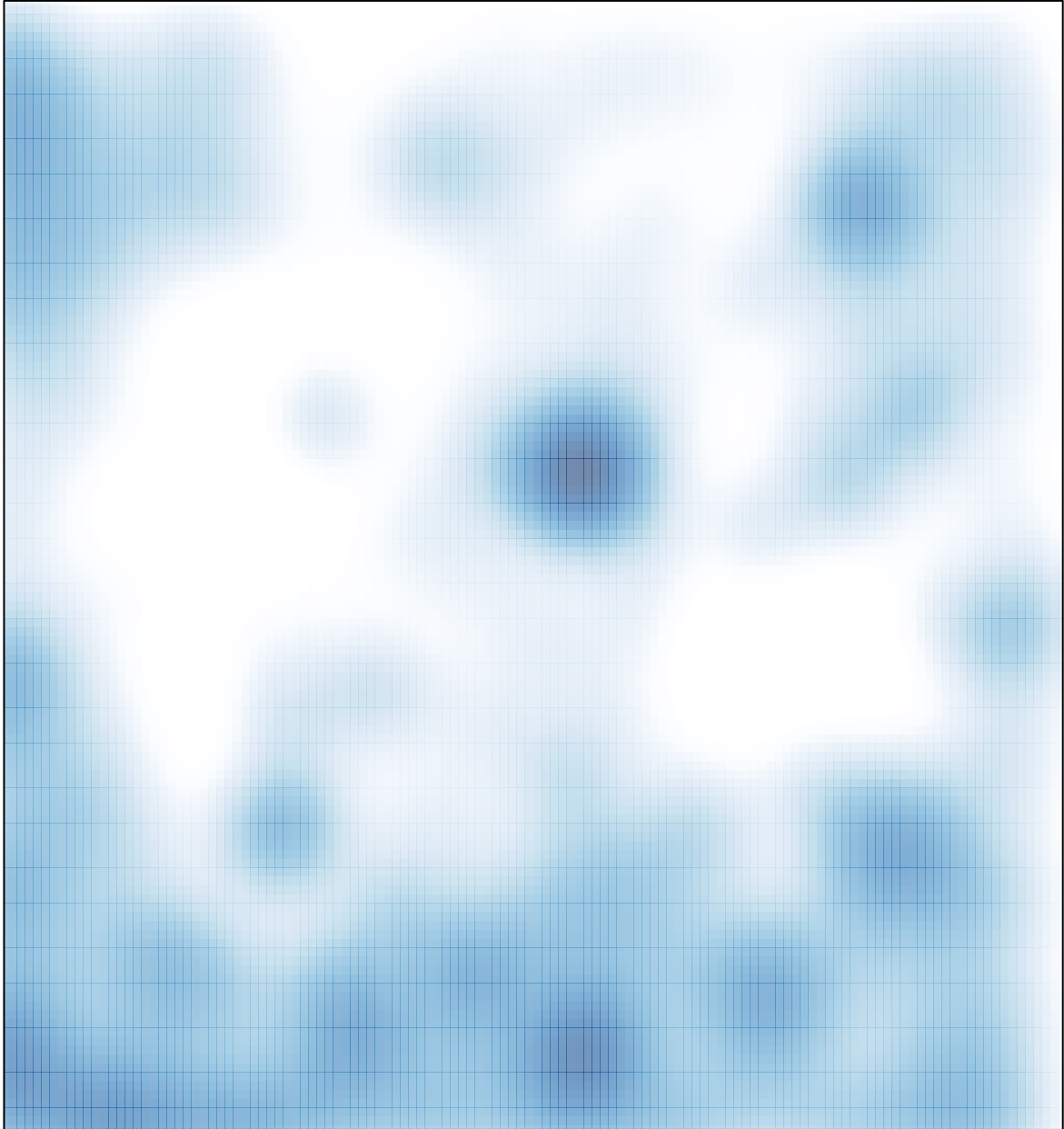


ion binding



features = 4992
chi-square p = 0.13

ion binding



features = 4992 , max = 41