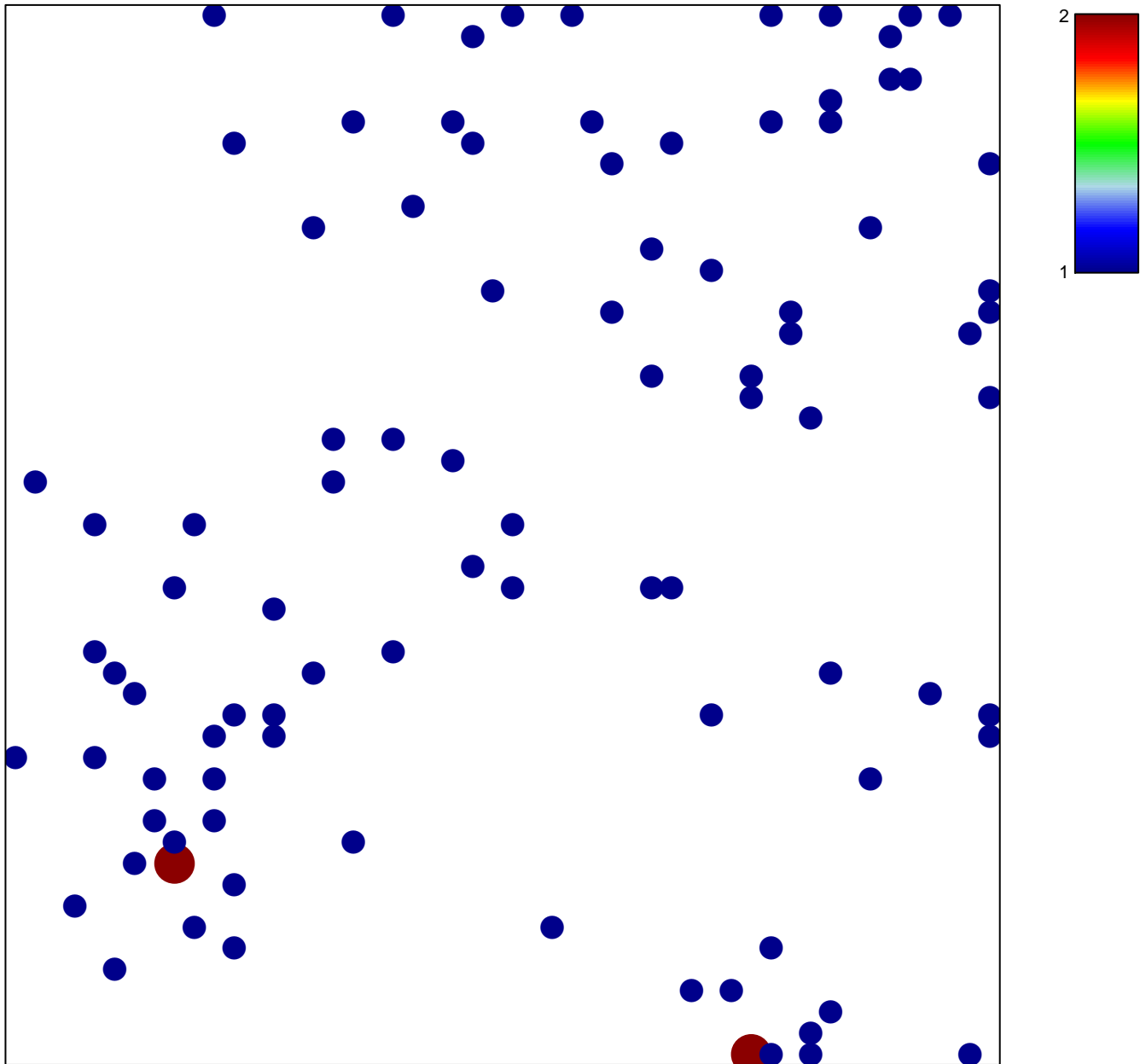
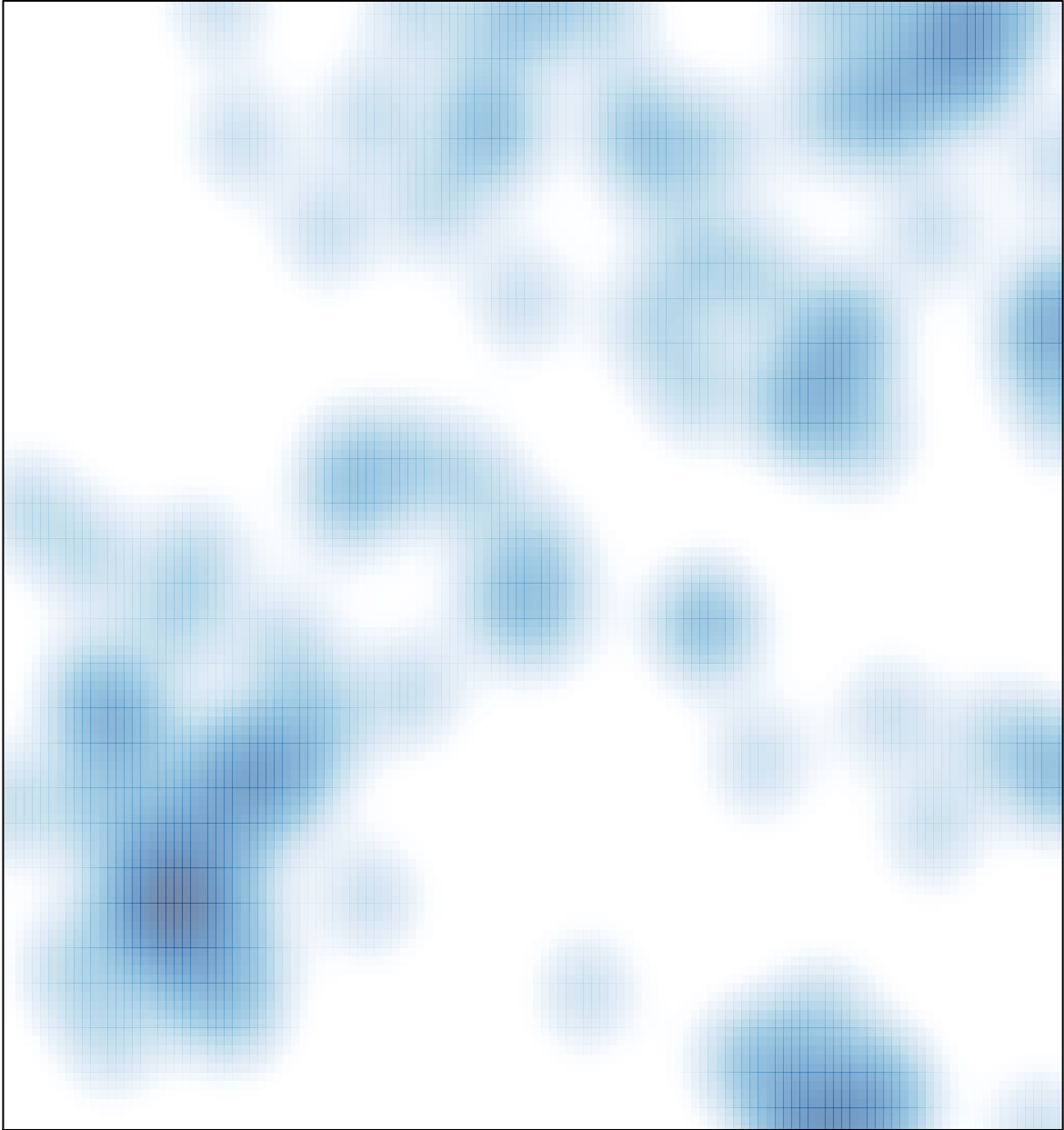


chromatin modification



features = 96
chi-square p = 0.84

chromatin modification



features = 96 , max = 2