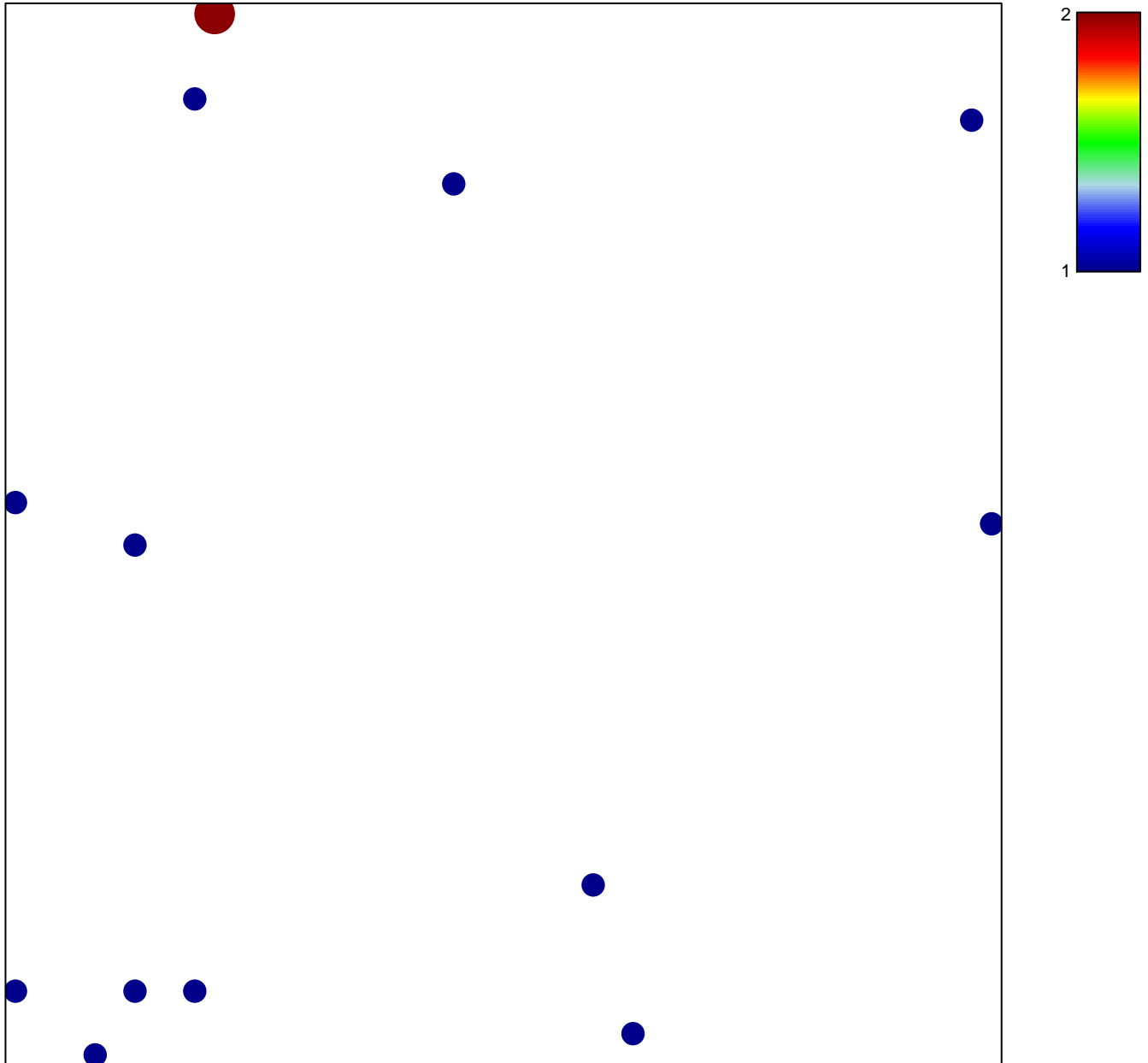
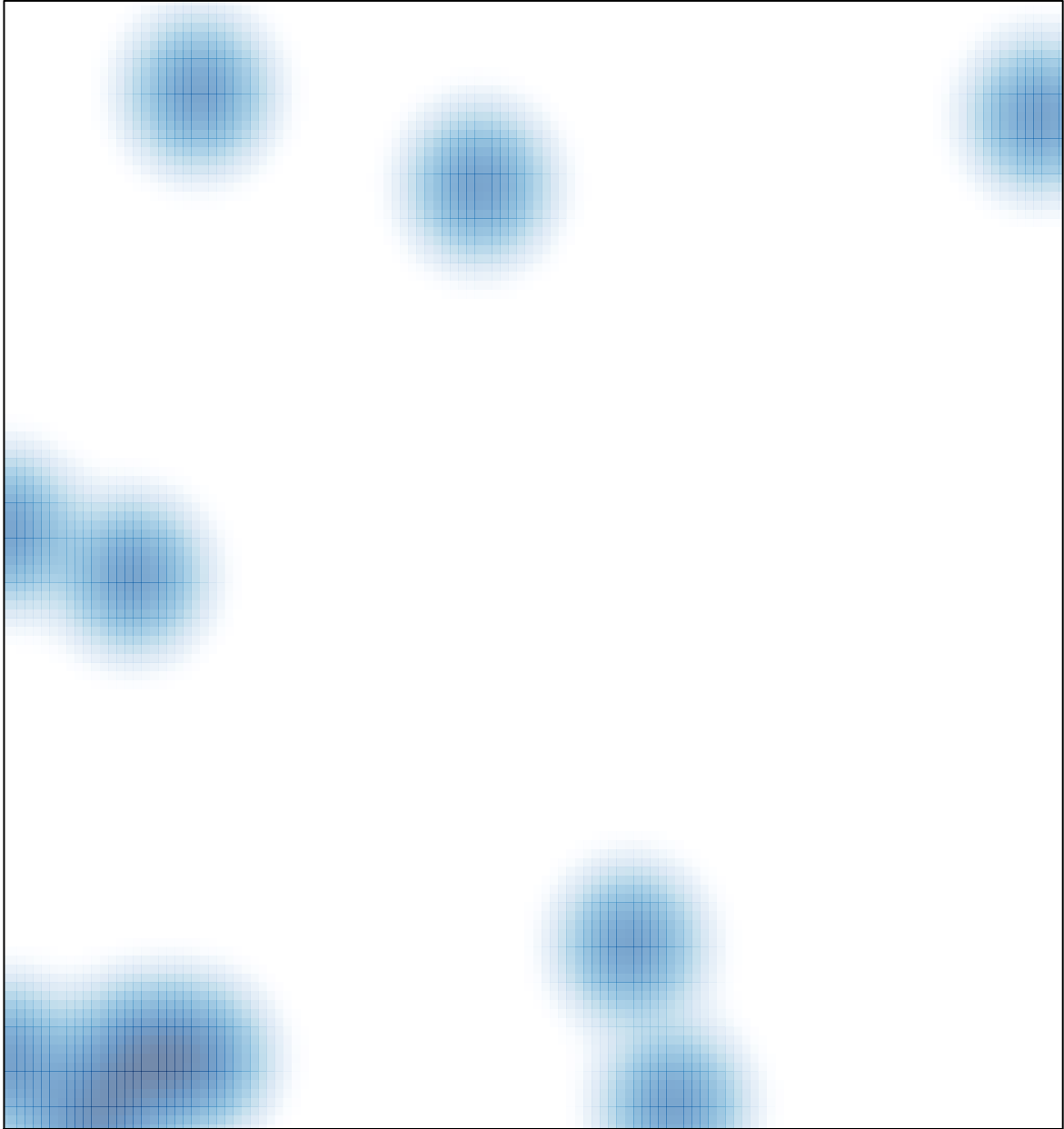


nucleosome disassembly



features = 14
chi-square p = 0.82

nucleosome disassembly



features = 14 , max = 2