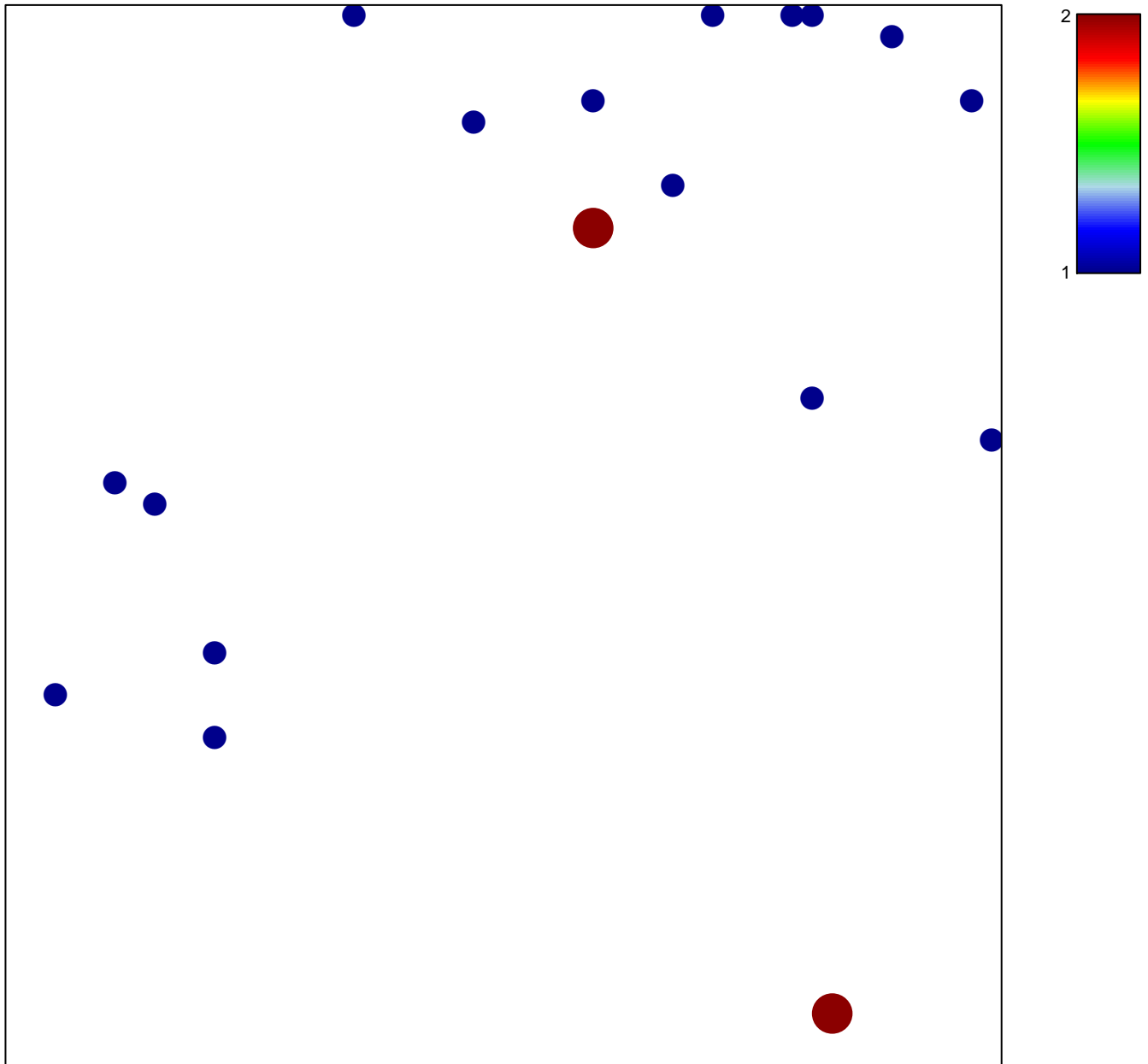


spliceosomal complex assembly



features = 20
chi-square p = 0.82

spliceosomal complex assembly



features = 20 , max = 2