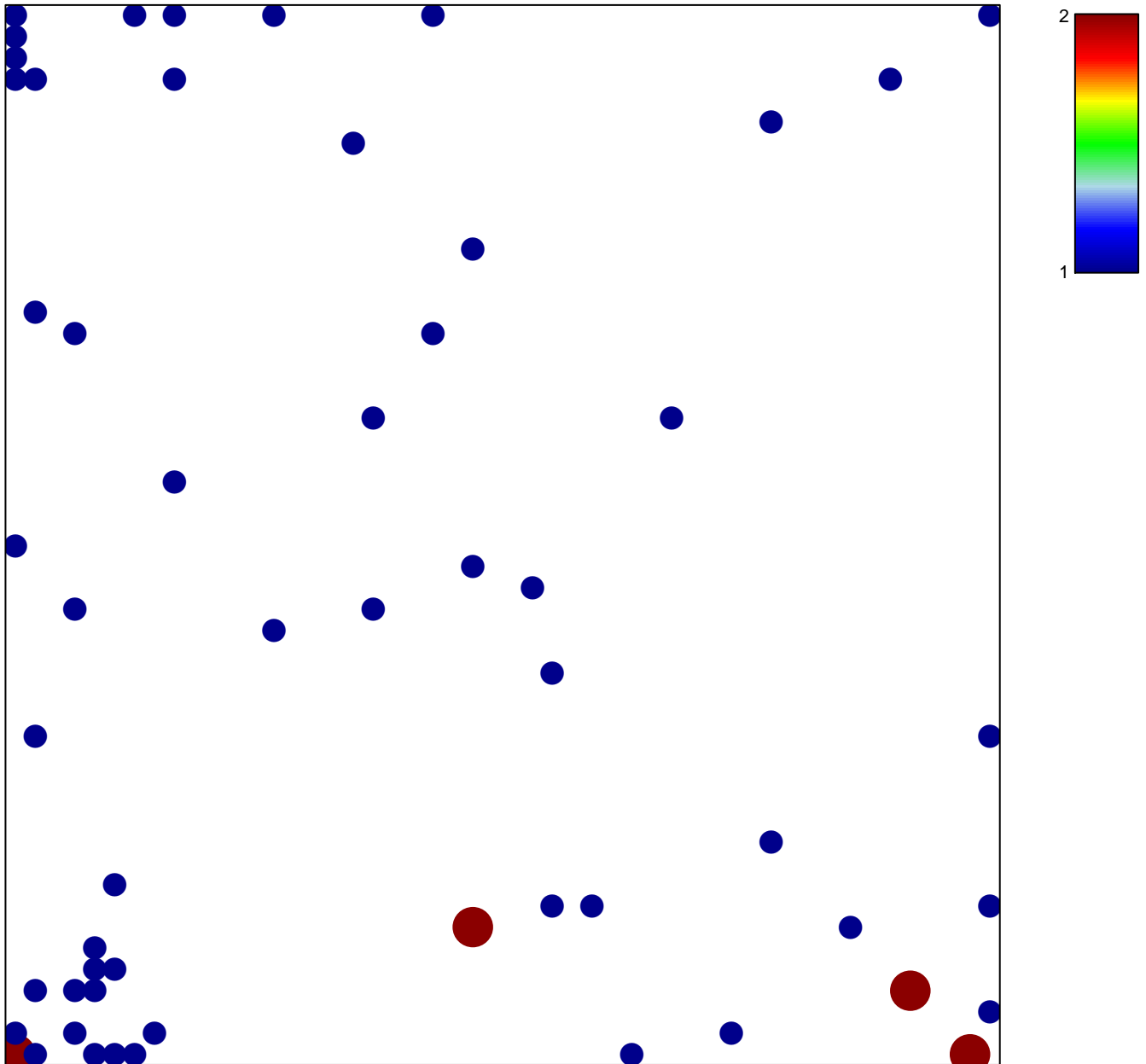
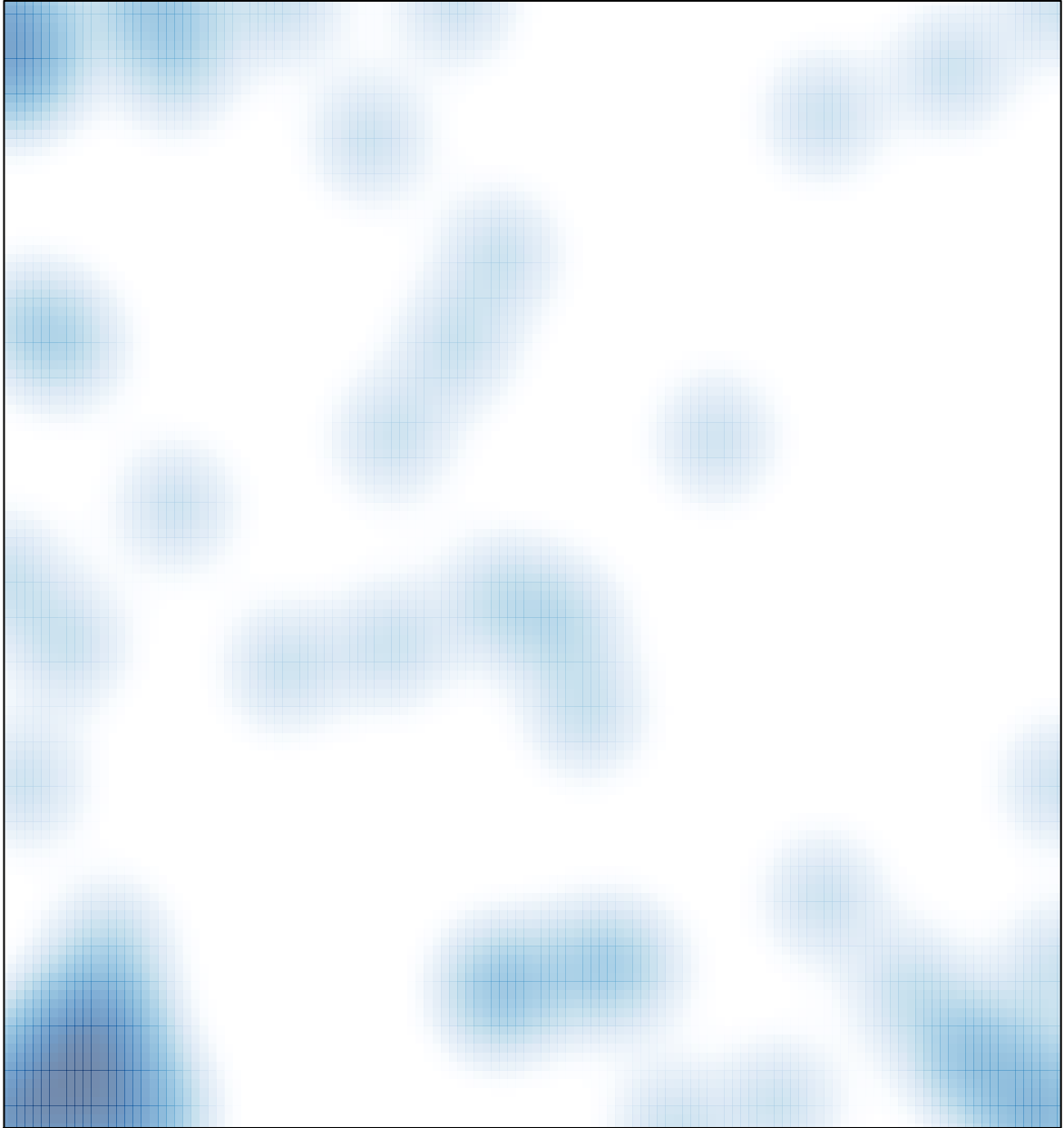


Christensen_hypomethylated_in_secondary_glioblastoma



features = 60
chi-square p = 0.75

ristensen_hypomethylated_in_secondary_glioblasto



features = 60 , max = 2