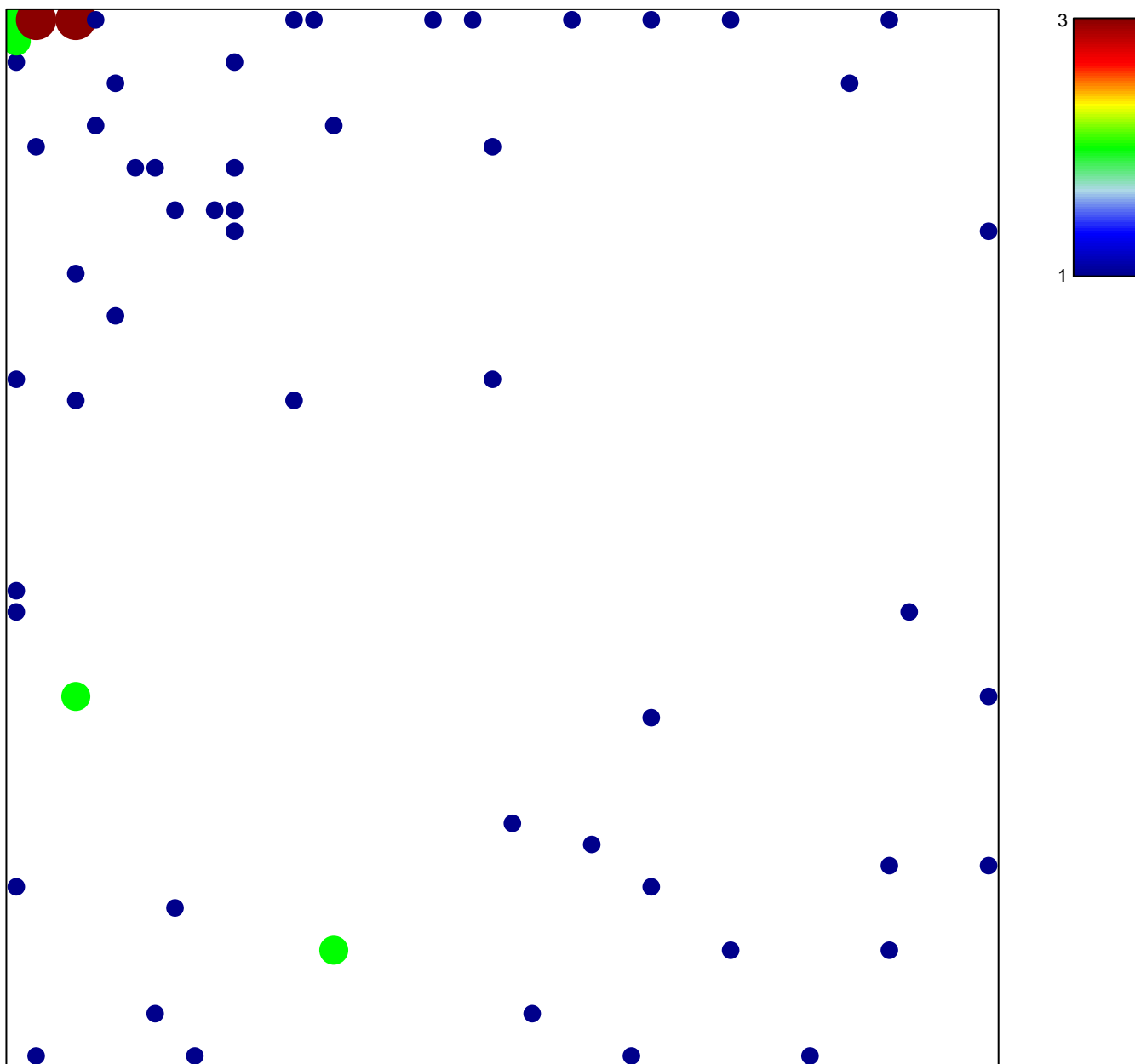
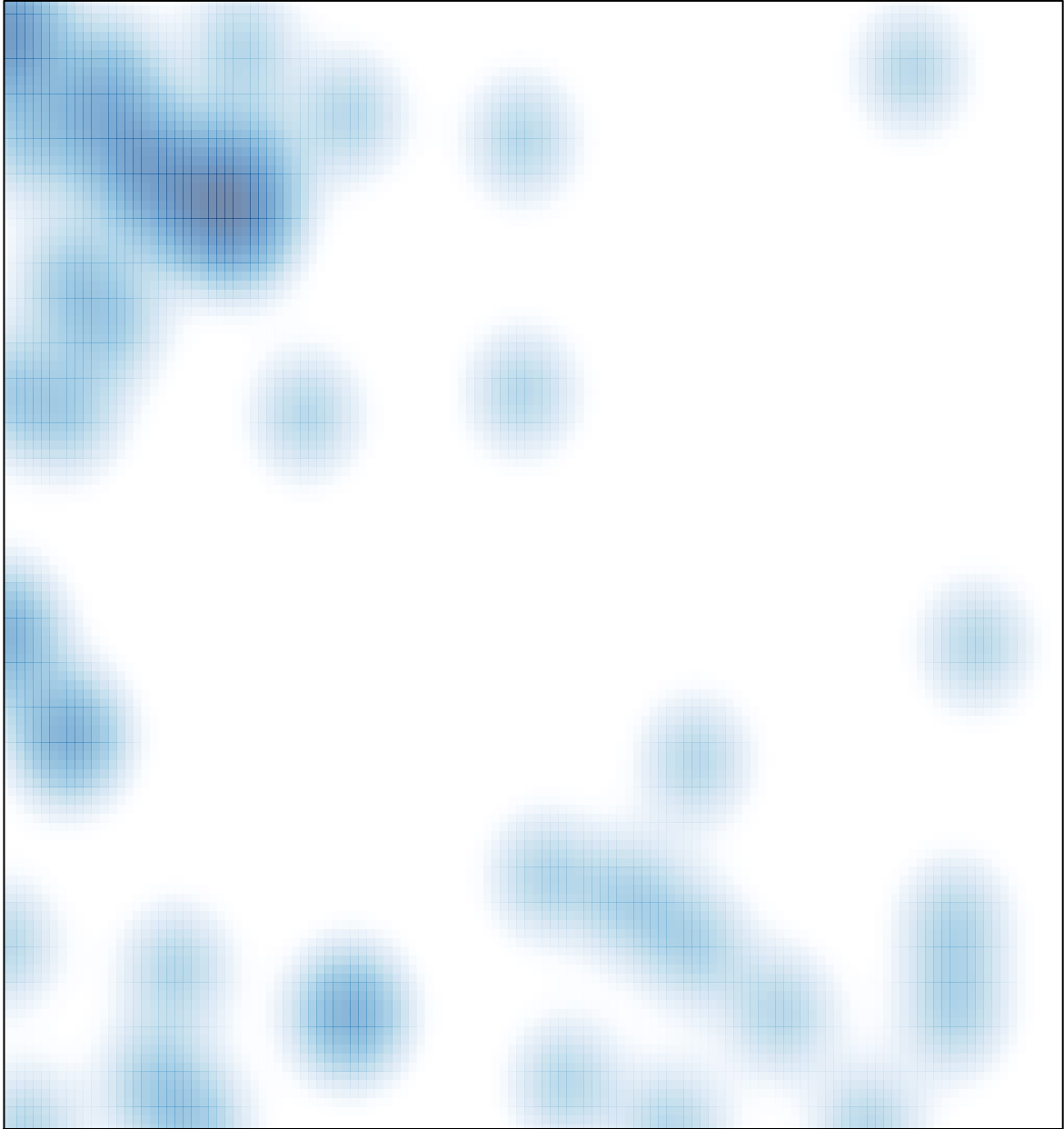


double-strand break repair via homologous recombination



features = 65
chi-square p = 0.7

Double-strand break repair via homologous recombination



features = 65 , max = 3