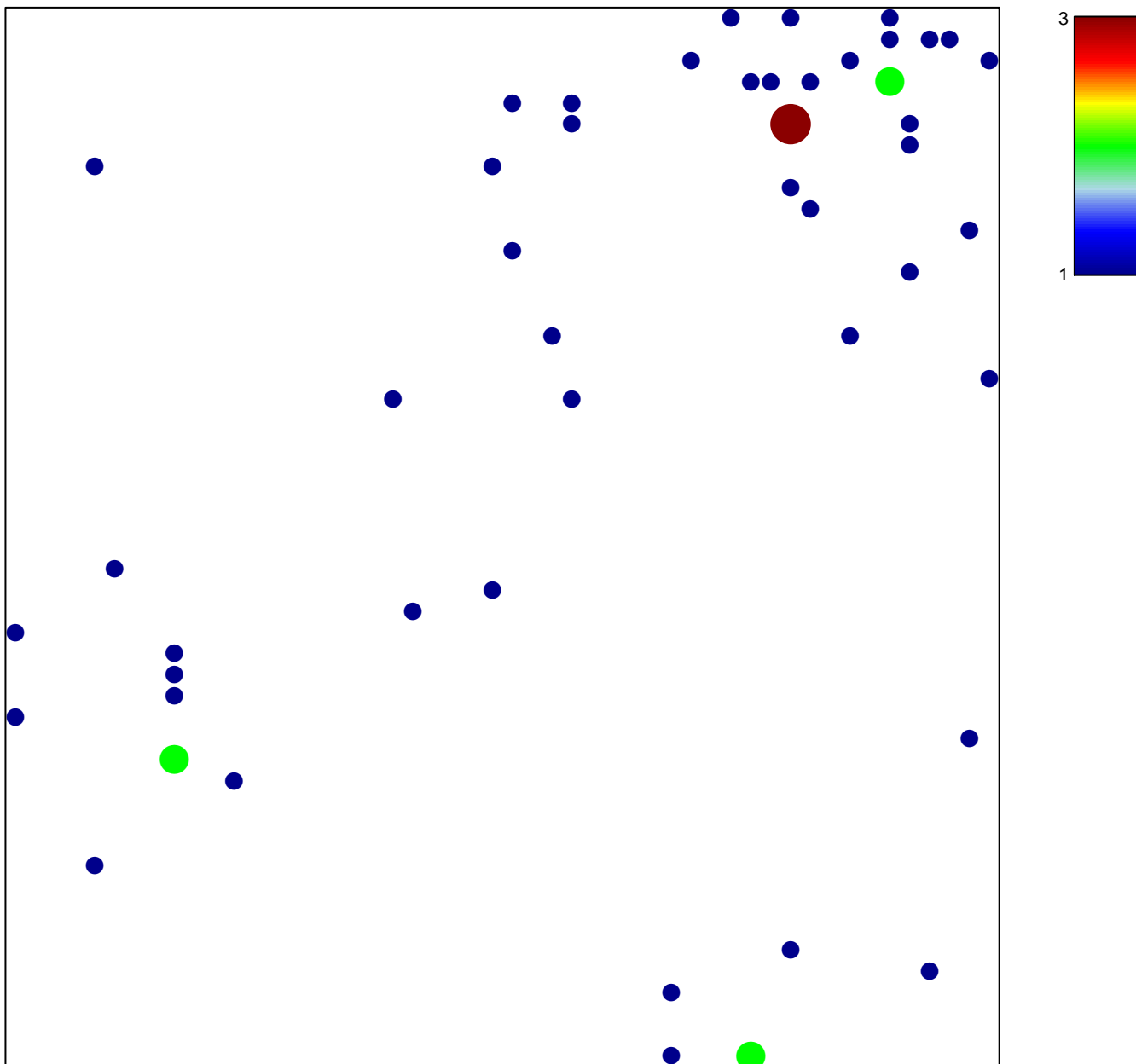
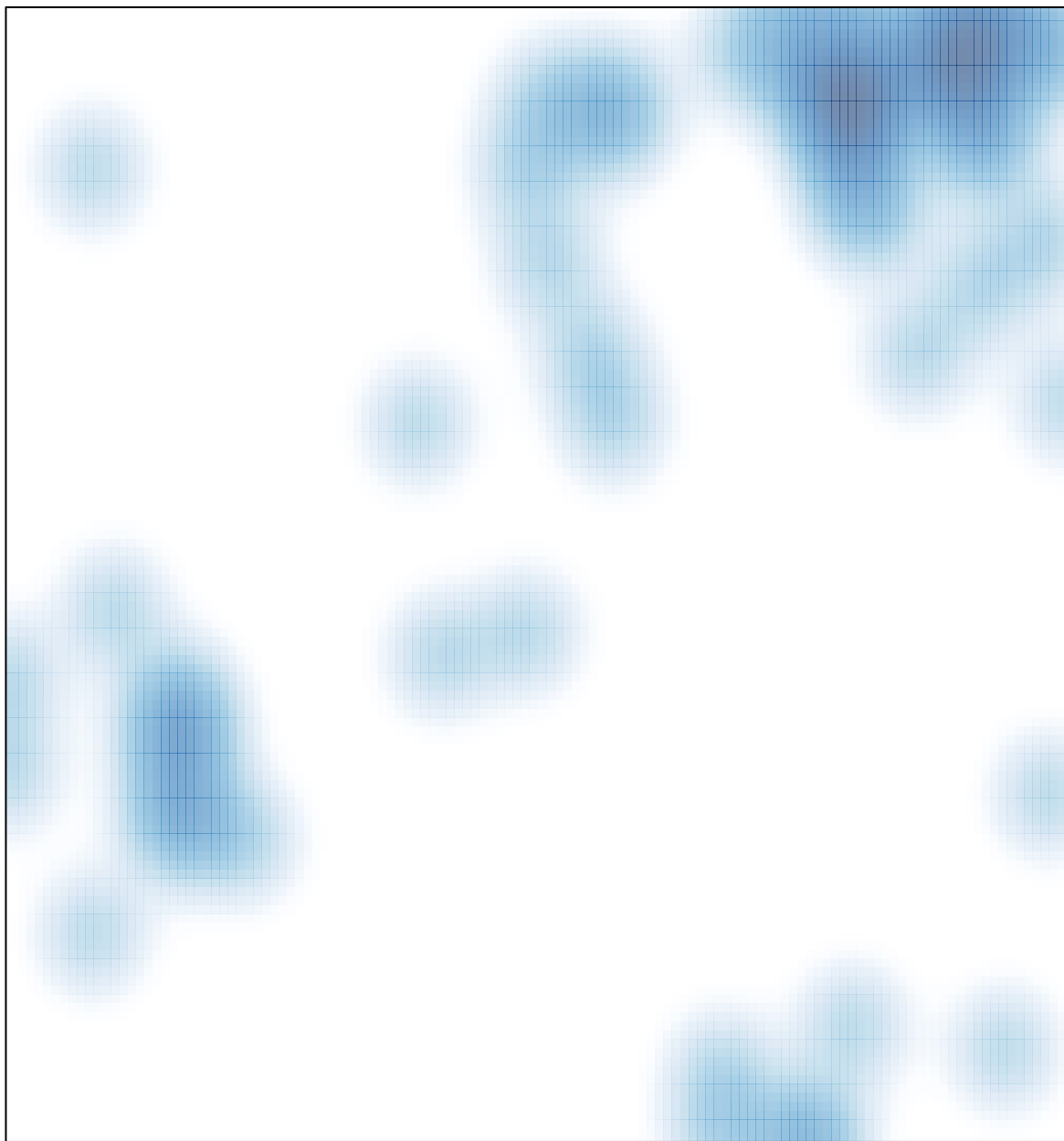


double-strand break repair via homologous recombination



features = 53
chi-square p = 0.8

Double-strand break repair via homologous recombination



features = 53 , max = 3