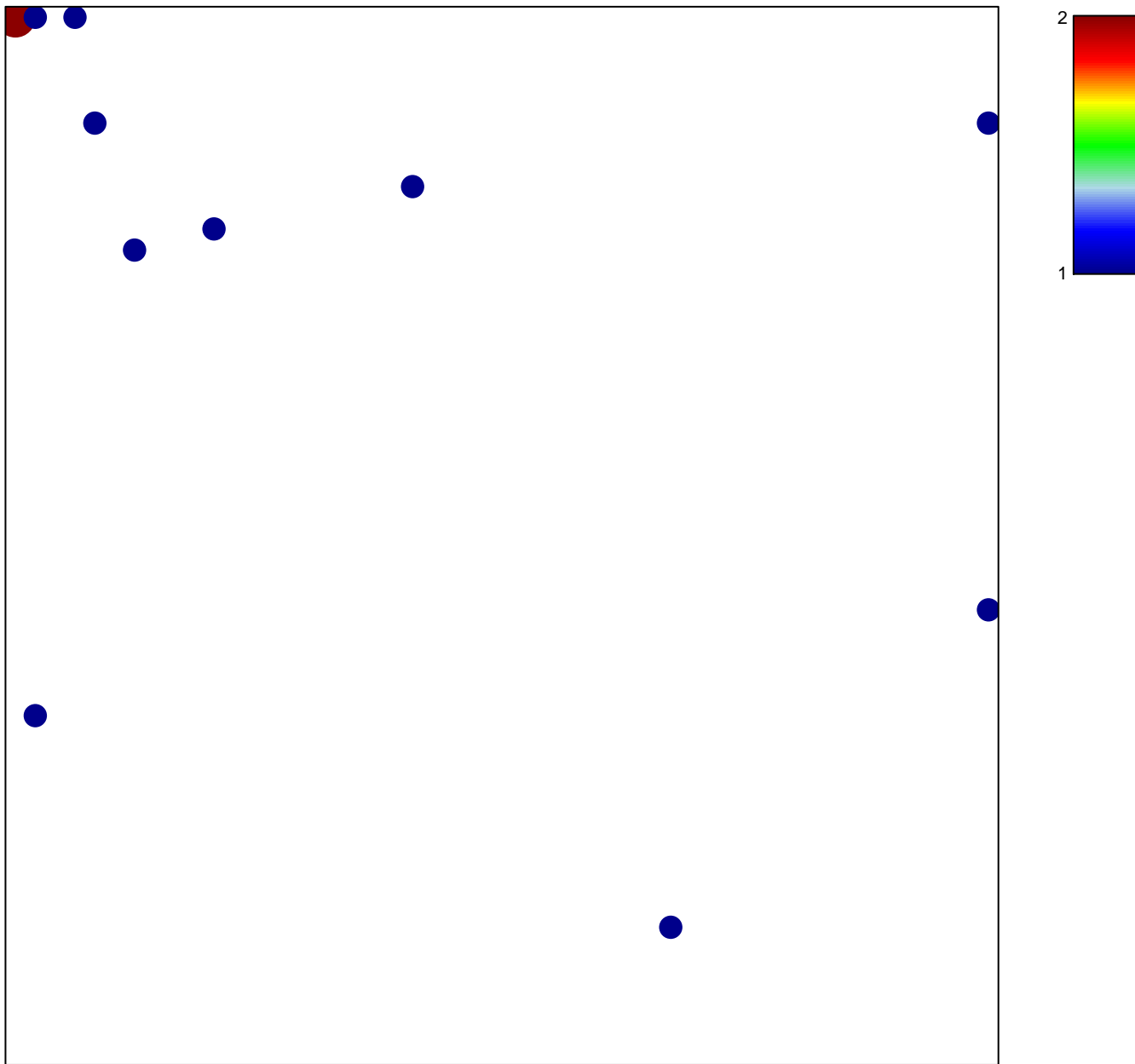


regulation of double-strand break repair via homologous recombination



features = 12
chi-square p = 0.8

h of double-strand break repair via homologous recombination



features = 12 , max = 2