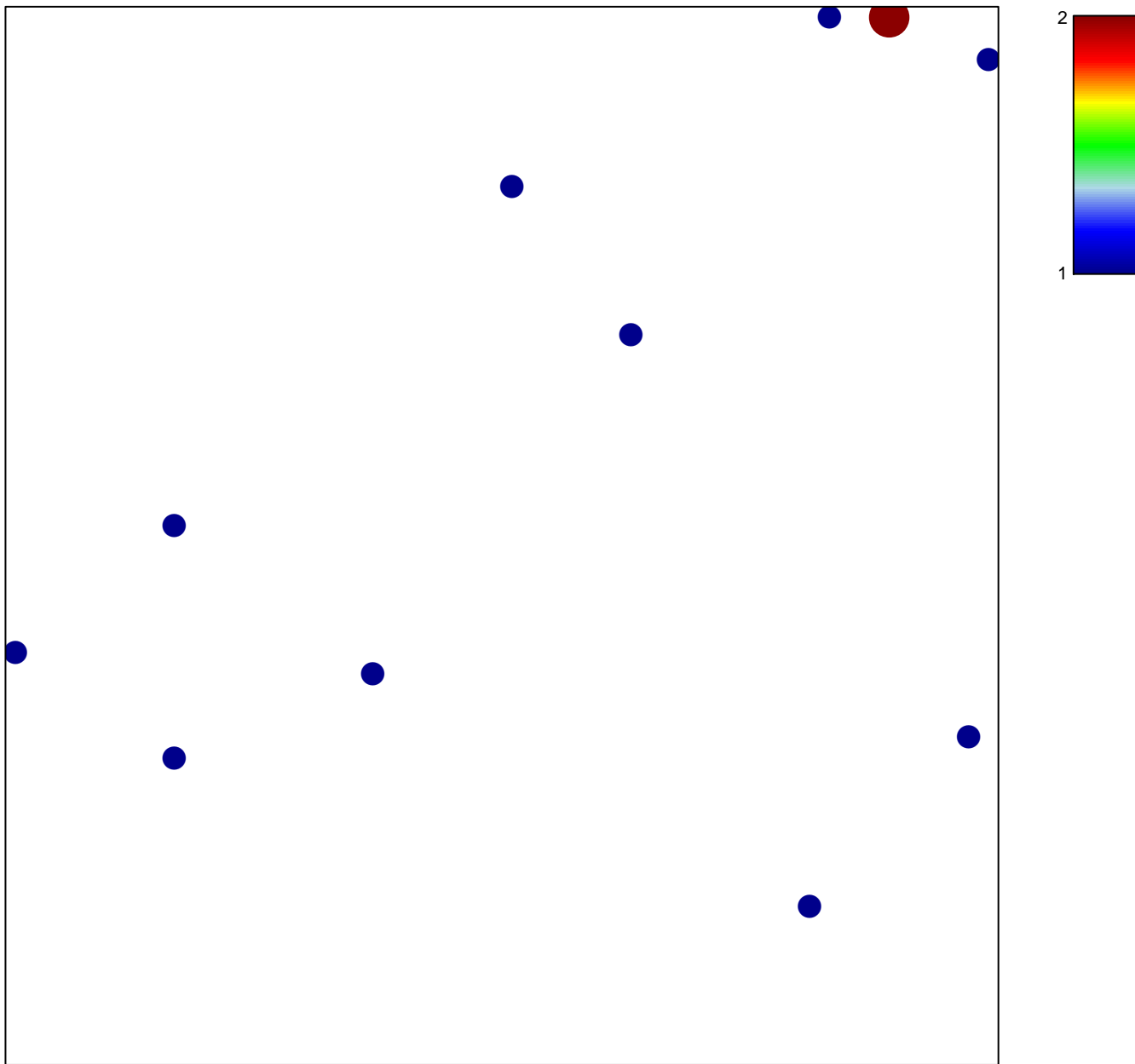


# regulation of double-strand break repair via homologous recombination



# features = 12  
chi-square p = 0.83

# h of double-strand break repair via homologous recombination



# features = 12 , max = 2