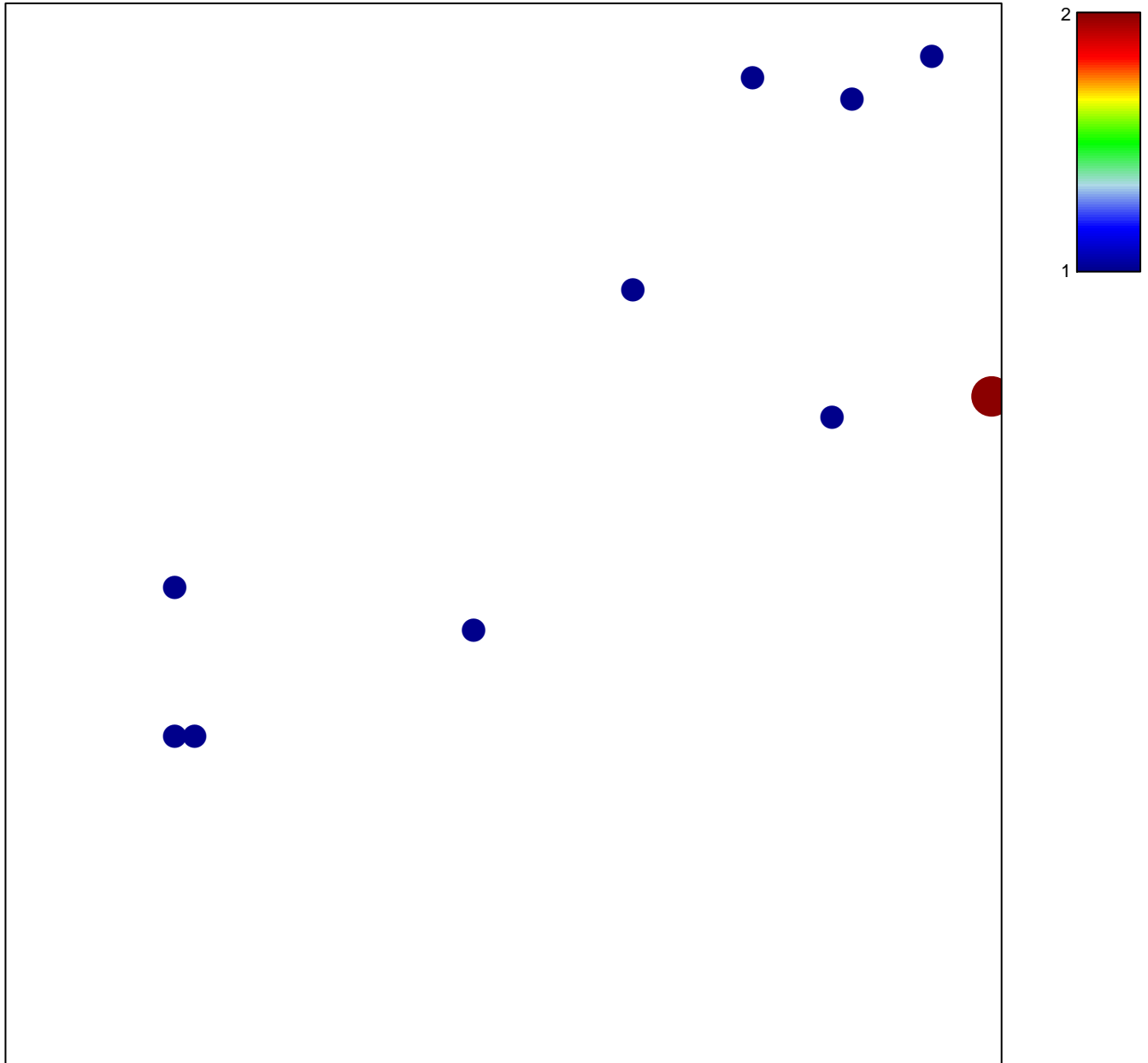


mismatched DNA binding



features = 11
chi-square p = 0.83

mismatched DNA binding



features = 11 , max = 2