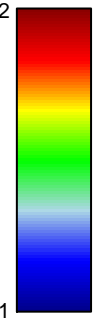
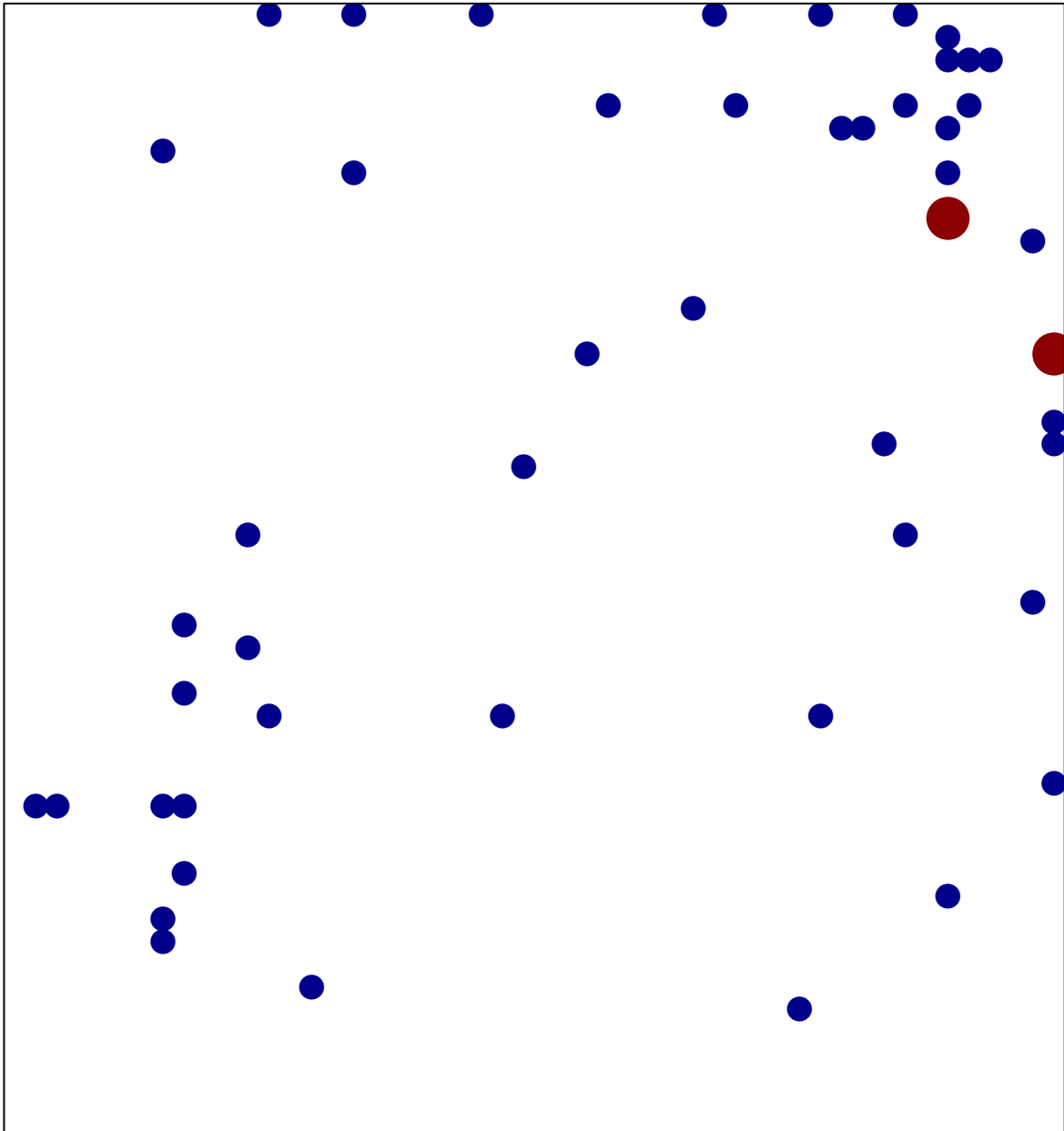


damaged DNA binding



features = 51
chi-square p = 0.79

damaged DNA binding



features = 51 , max = 2