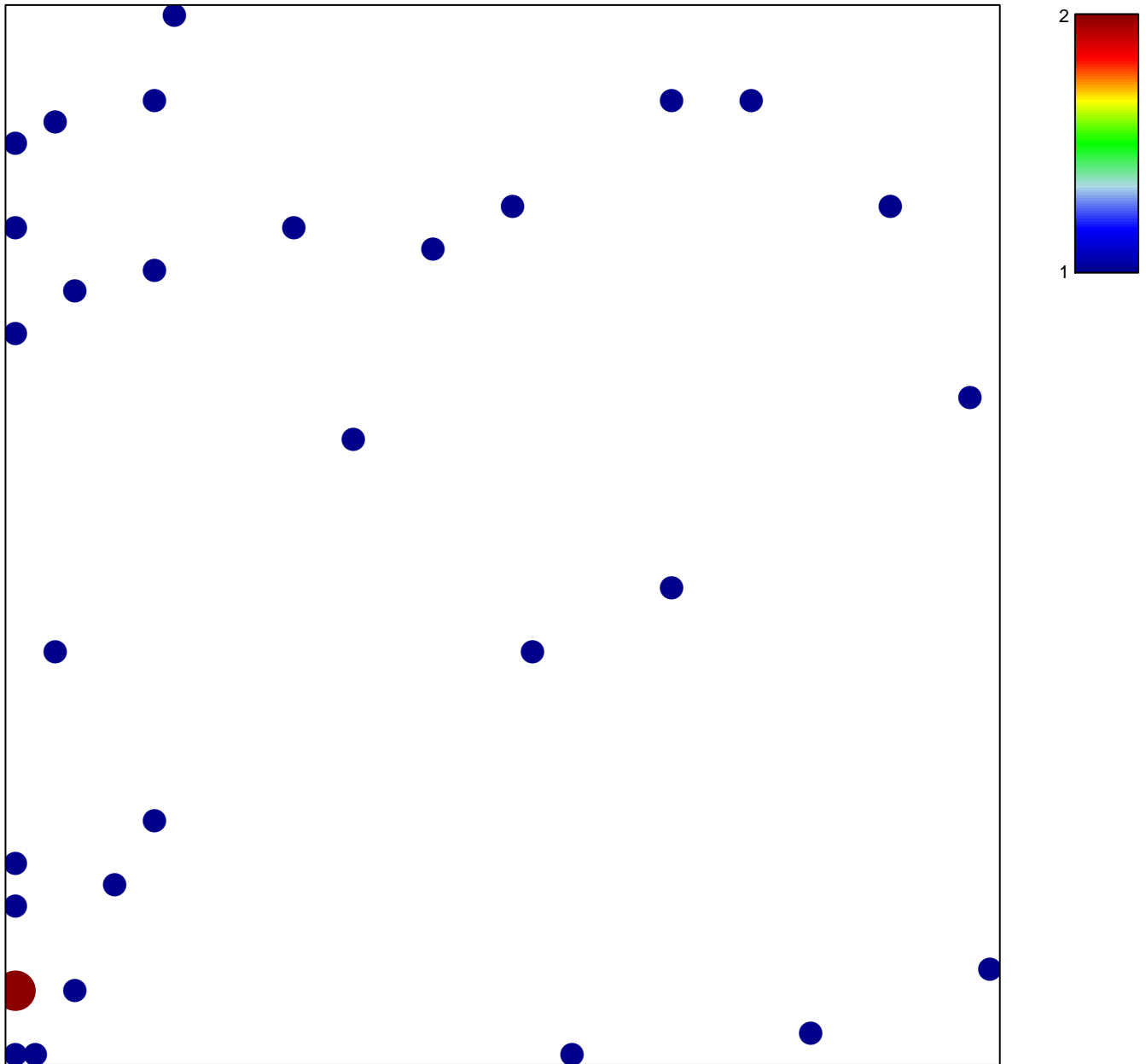


peptide binding



features = 31
chi-square p = 0.78

peptide binding



features = 31 , max = 2