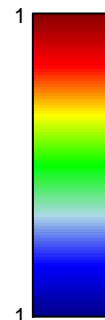
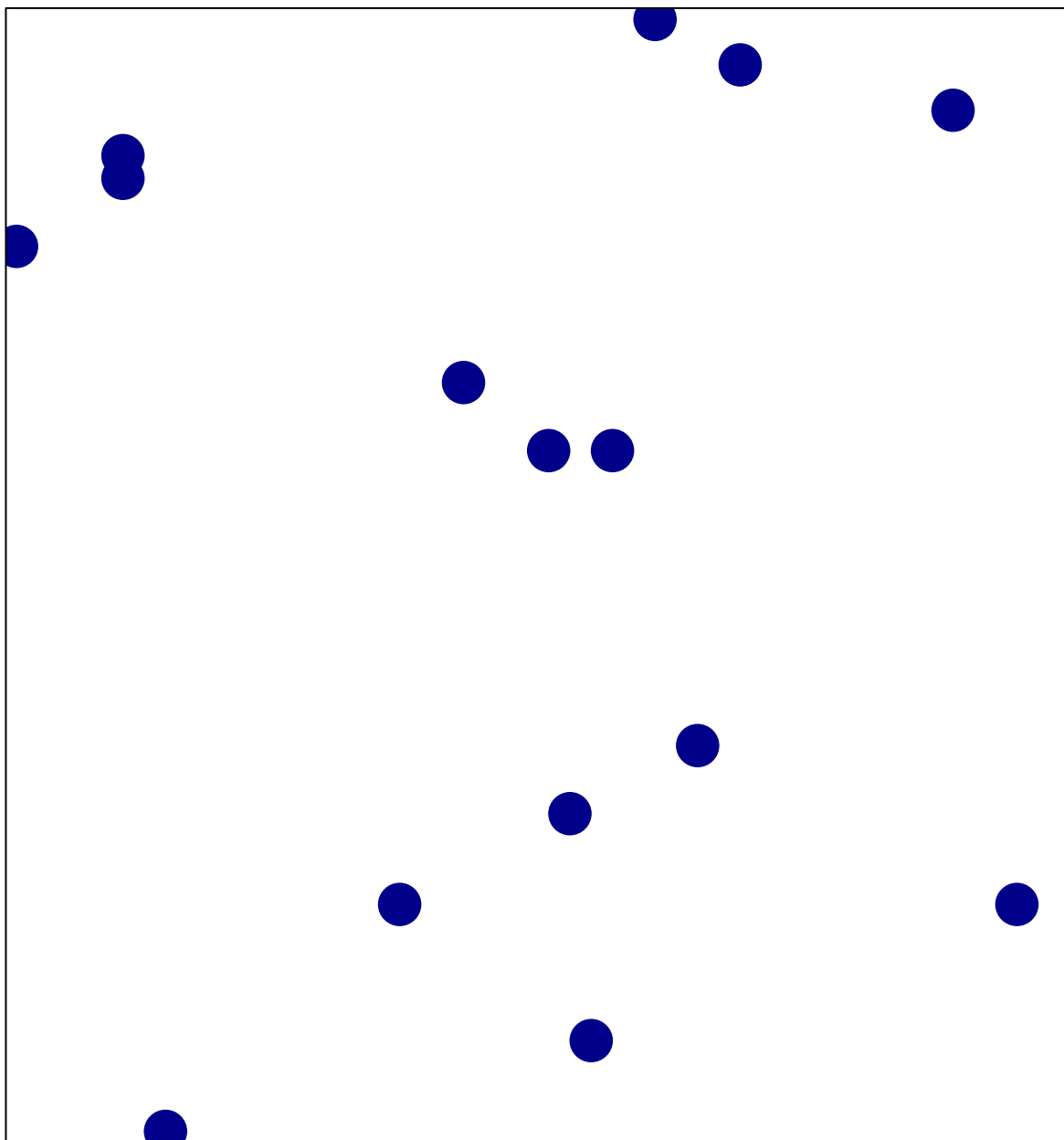


amino acid binding



features = 15
chi-square p = 0.85

amino acid binding



features = 15 , max = 1