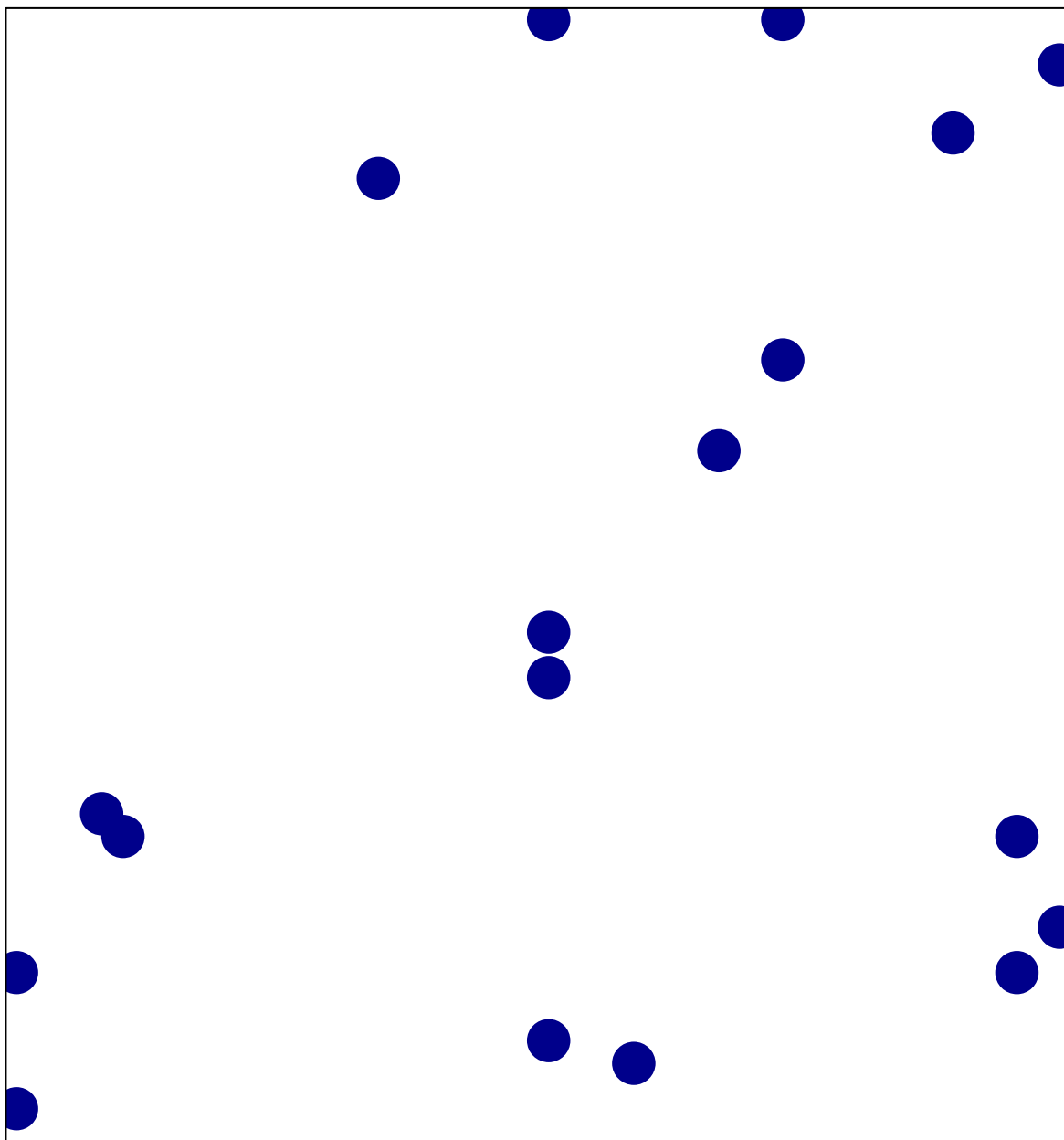
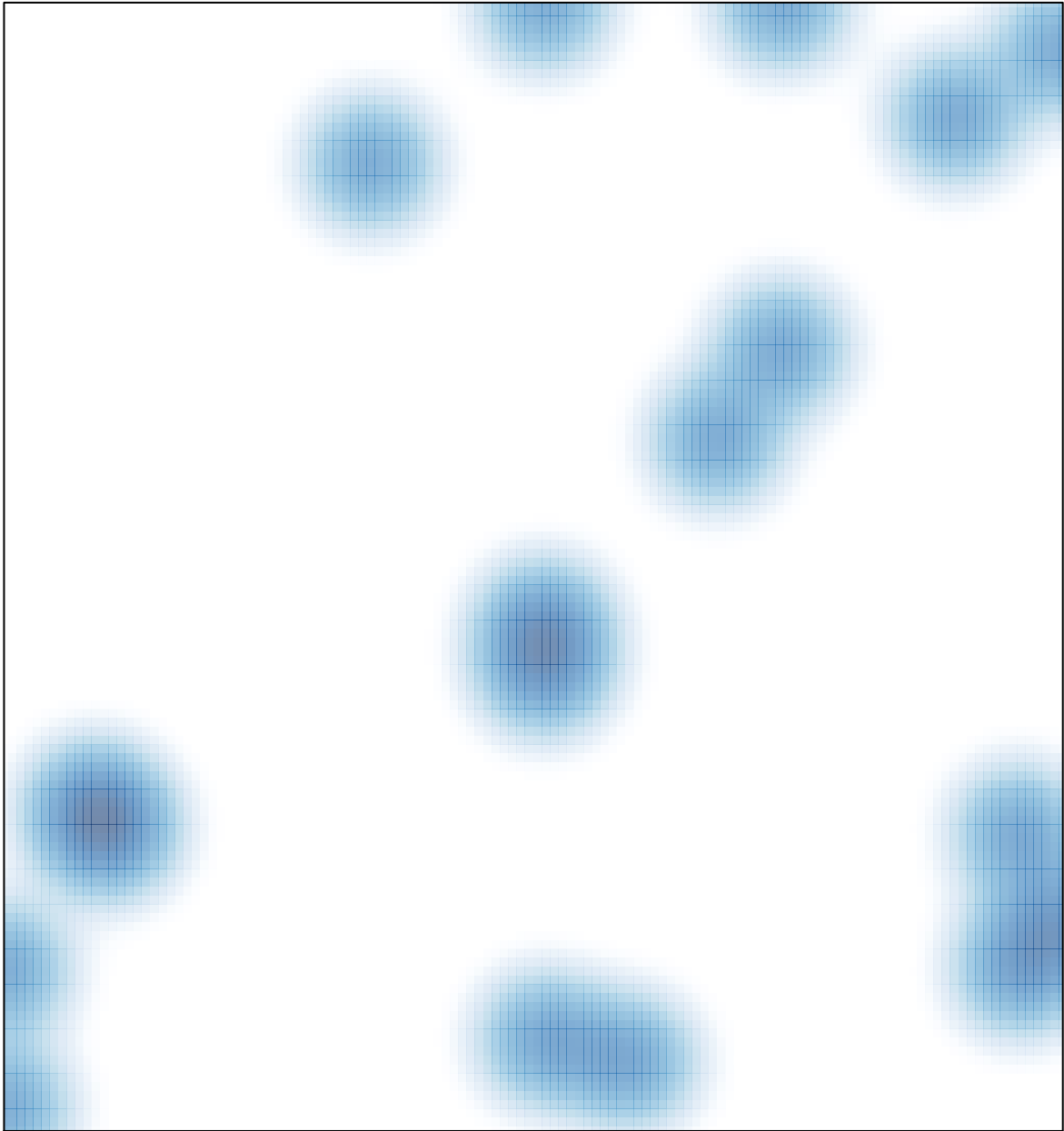


amino acid binding



features = 18
chi-square p = 0.82

amino acid binding



features = 18 , max = 1