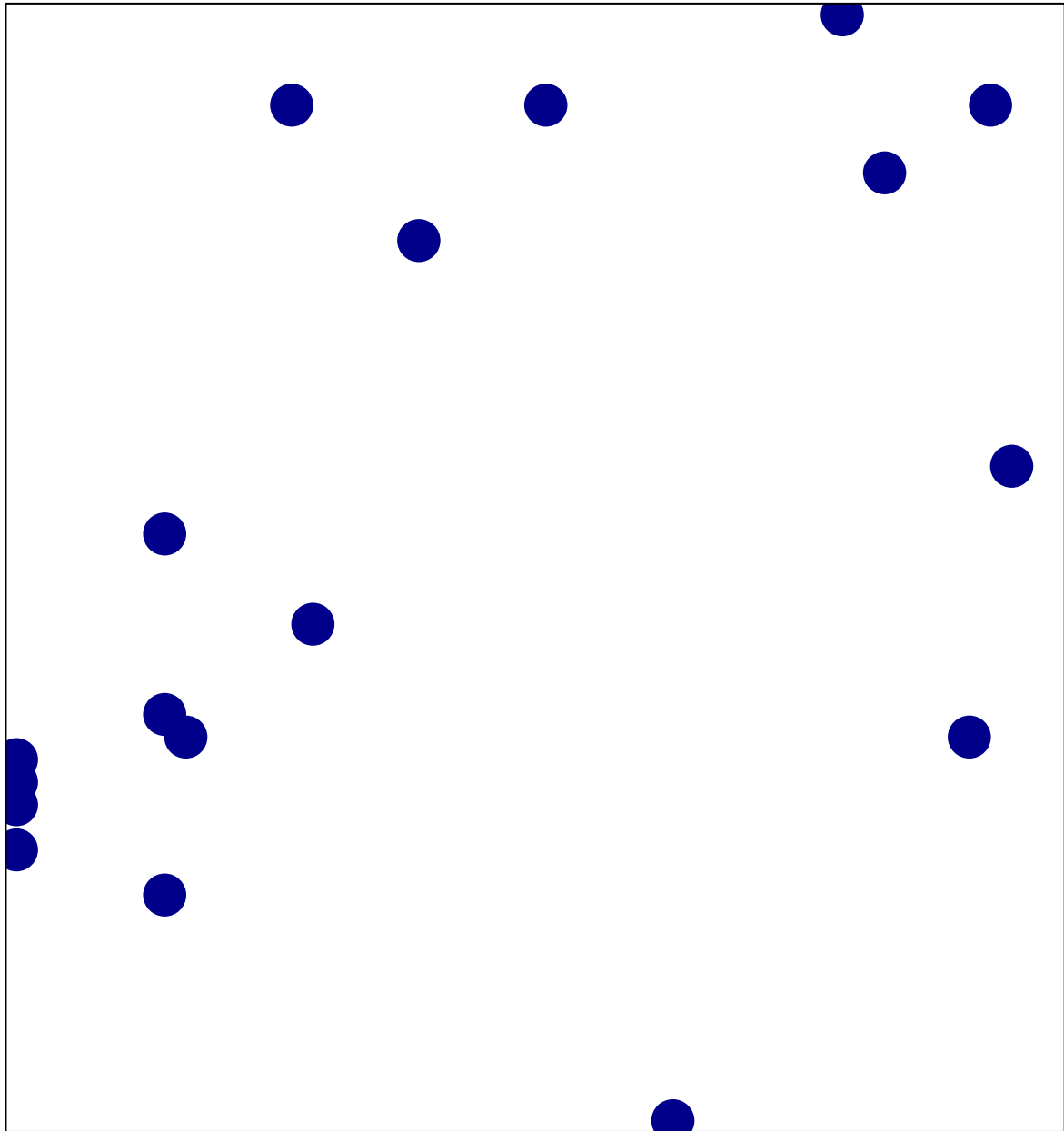
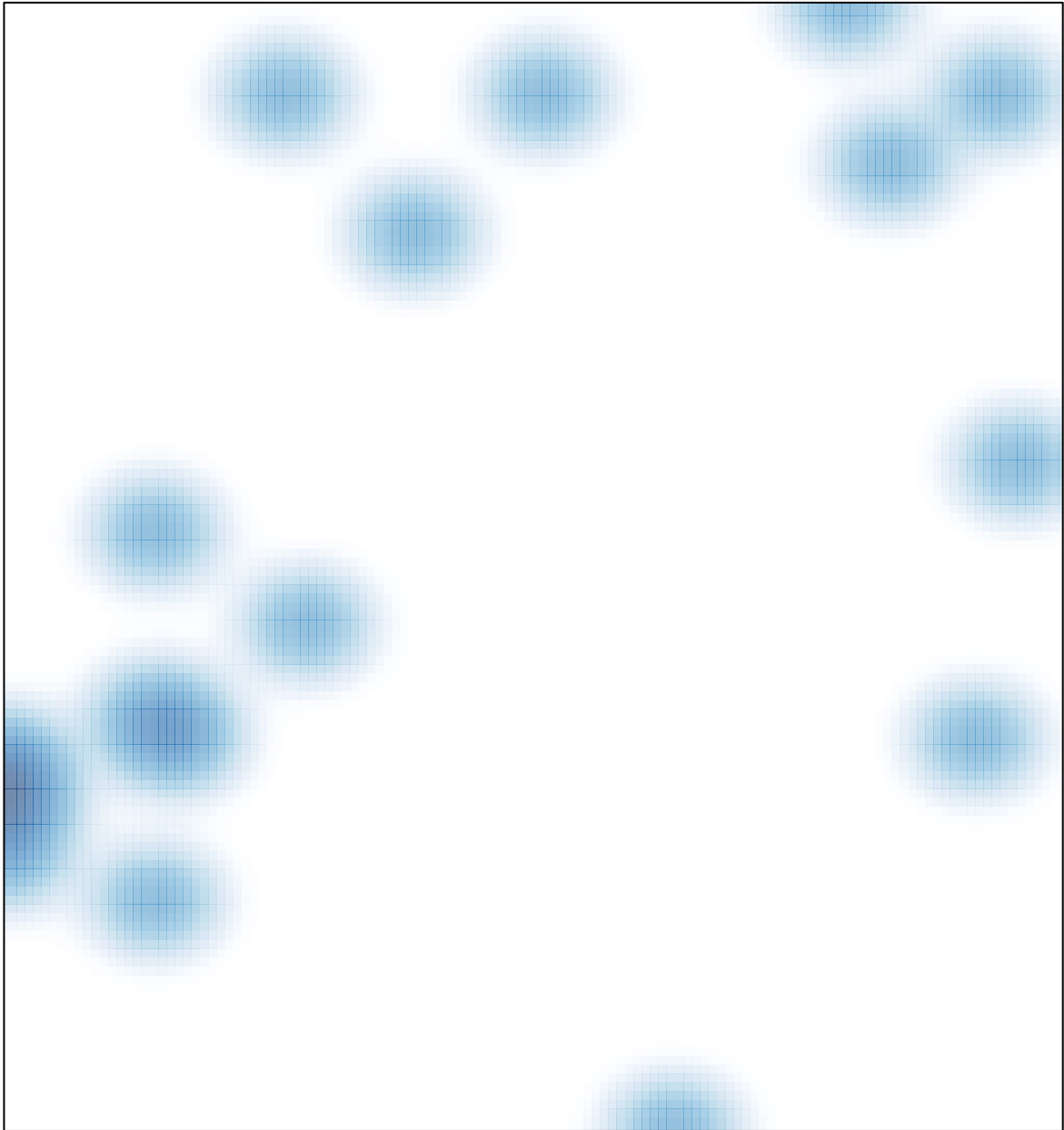


ribonucleoprotein complex binding



features = 18
chi-square p = 0.82

ribonucleoprotein complex binding



features = 18 , max = 1