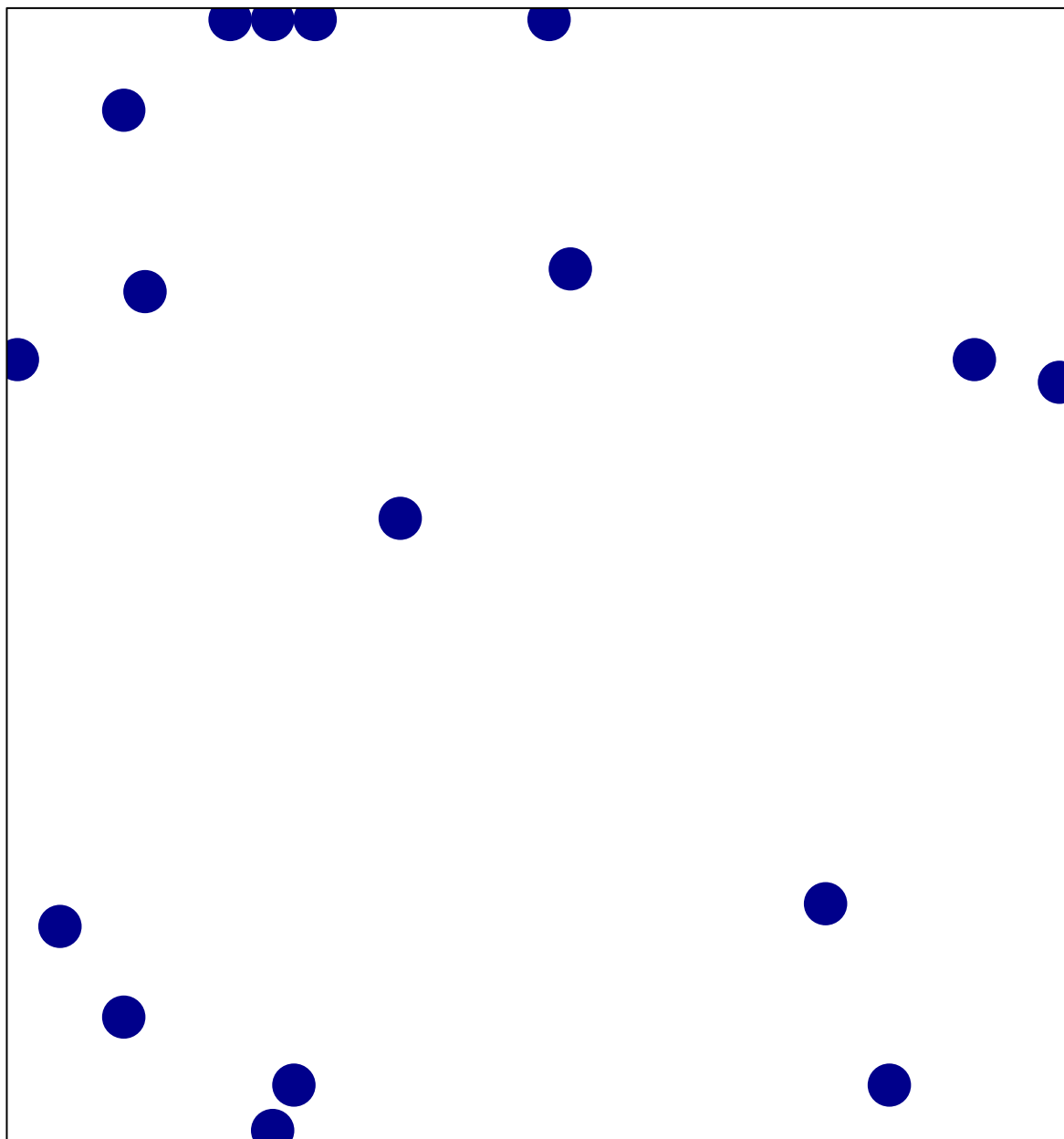
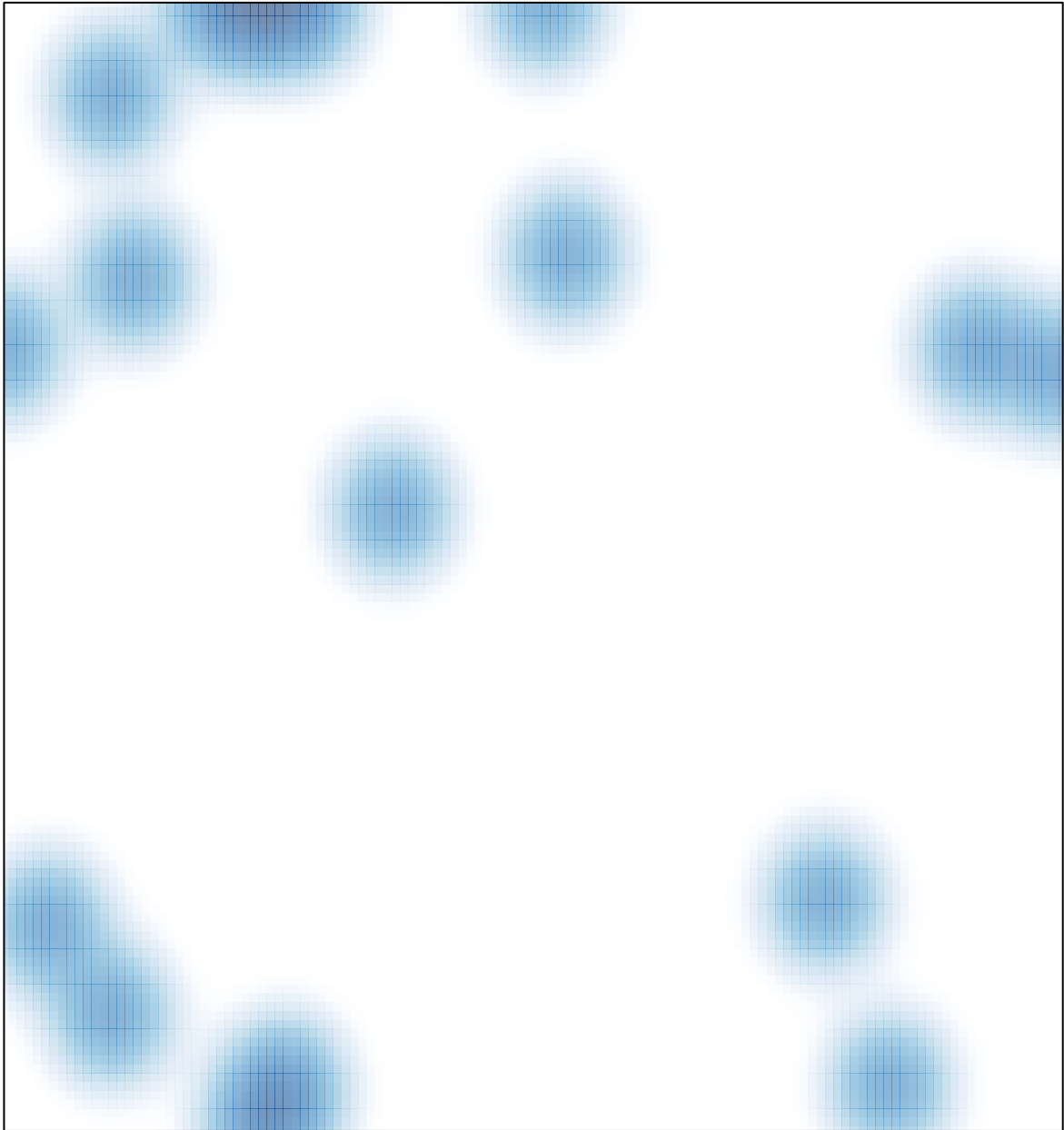


proline-rich region binding



features = 17
chi-square p = 0.82

proline-rich region binding



features = 17 , max = 1