proline−rich region binding
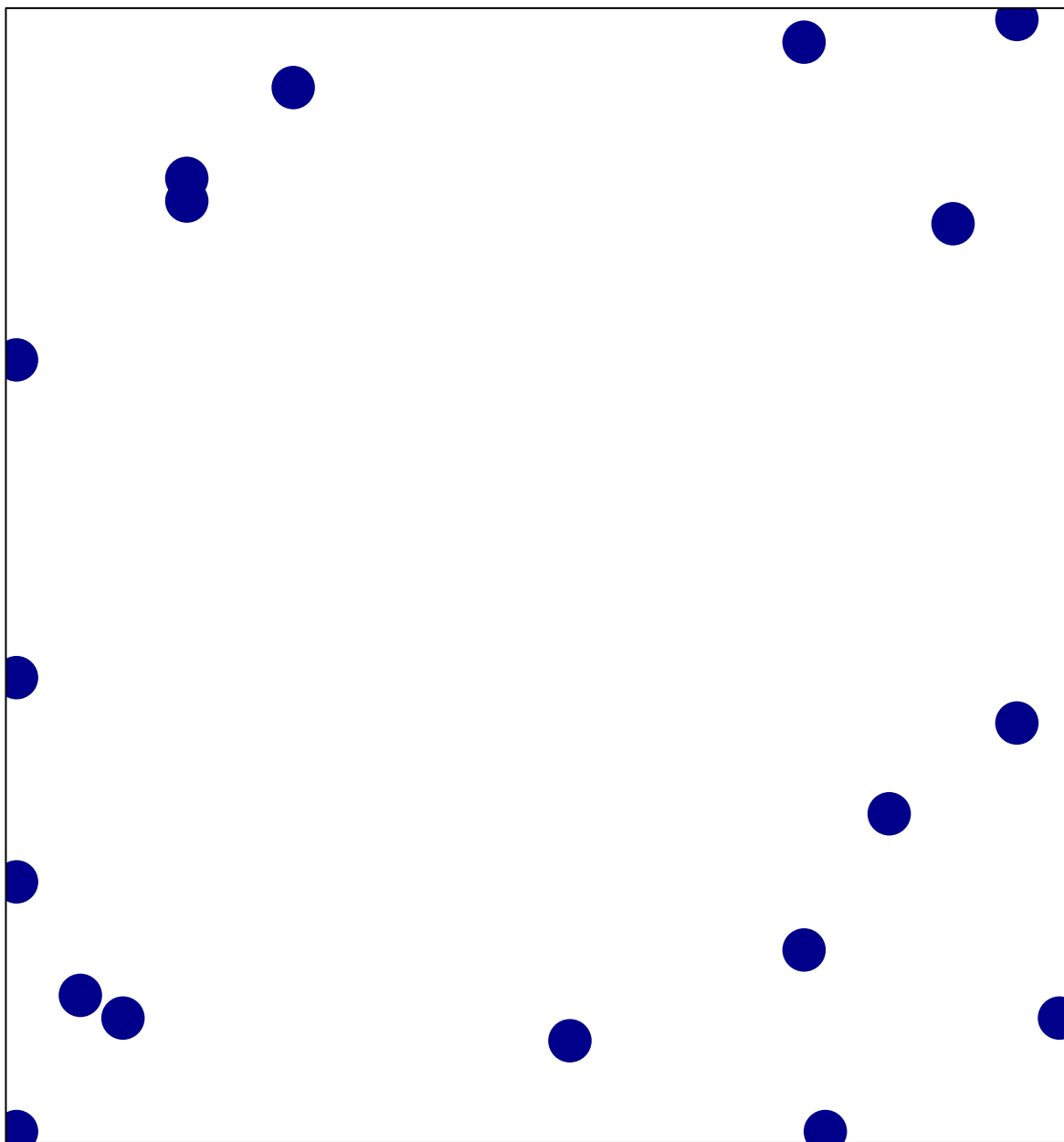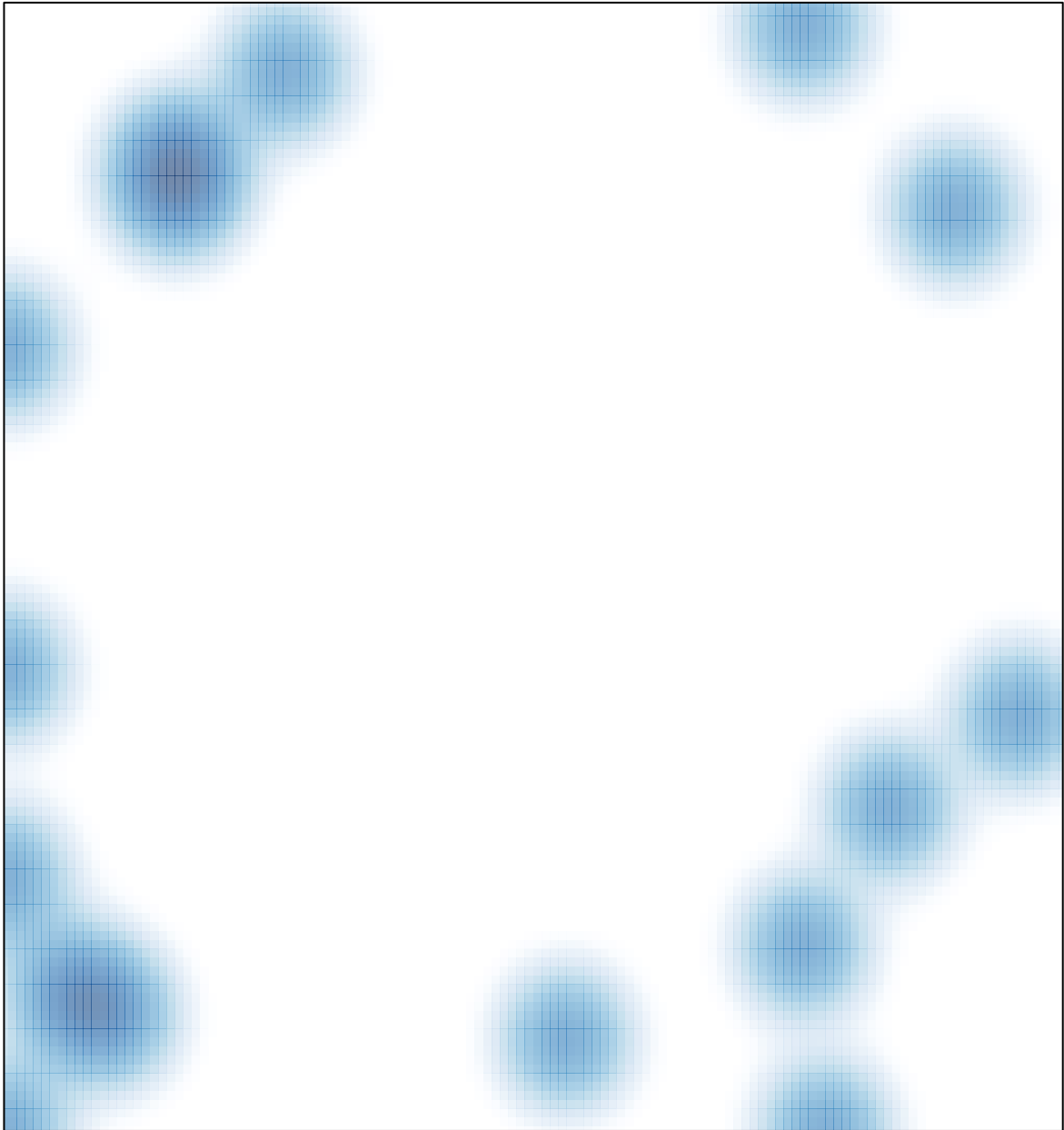

features = 18
chi−square p = 0.82

proline–rich region binding

features = 18 , max = 1