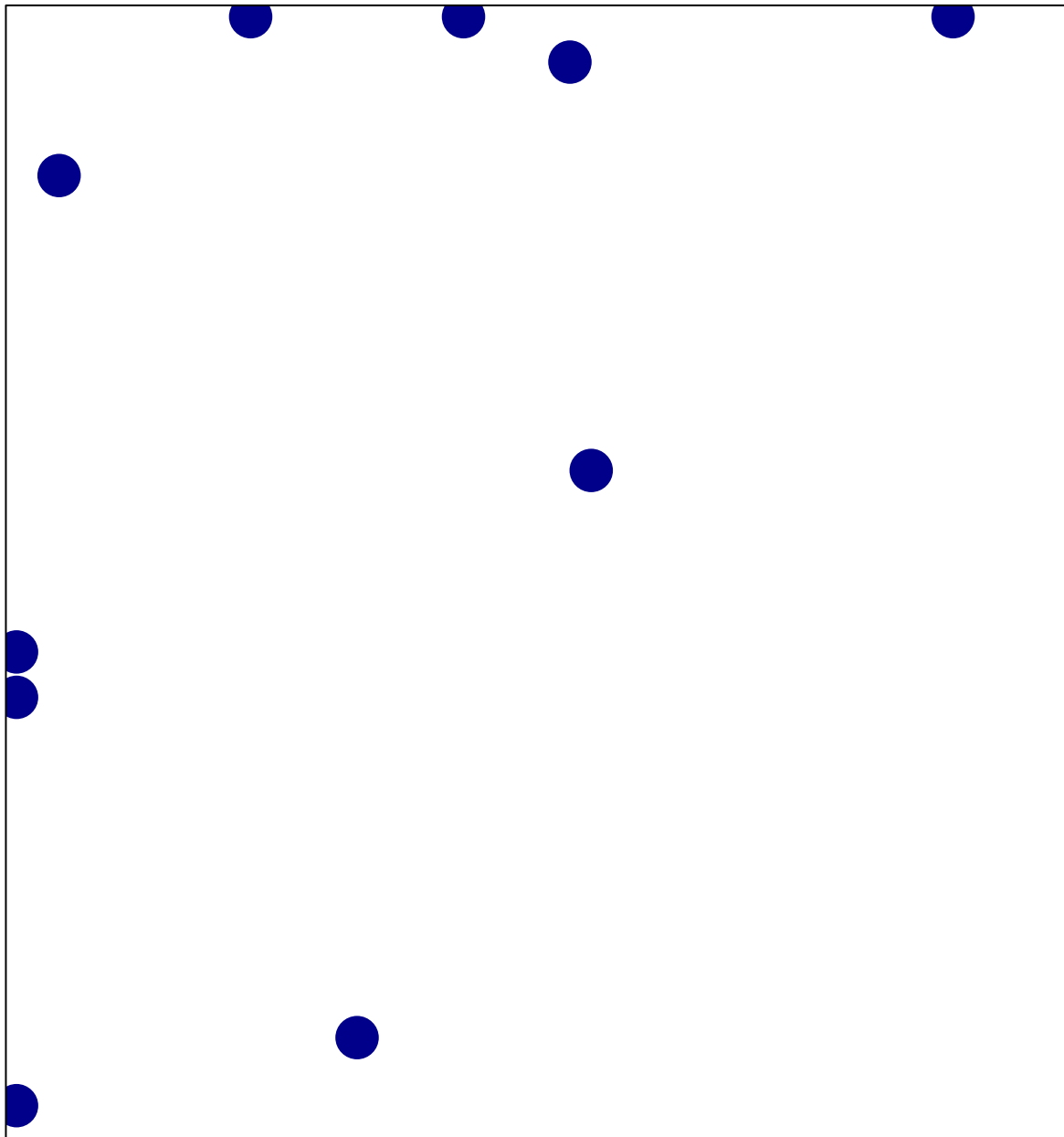
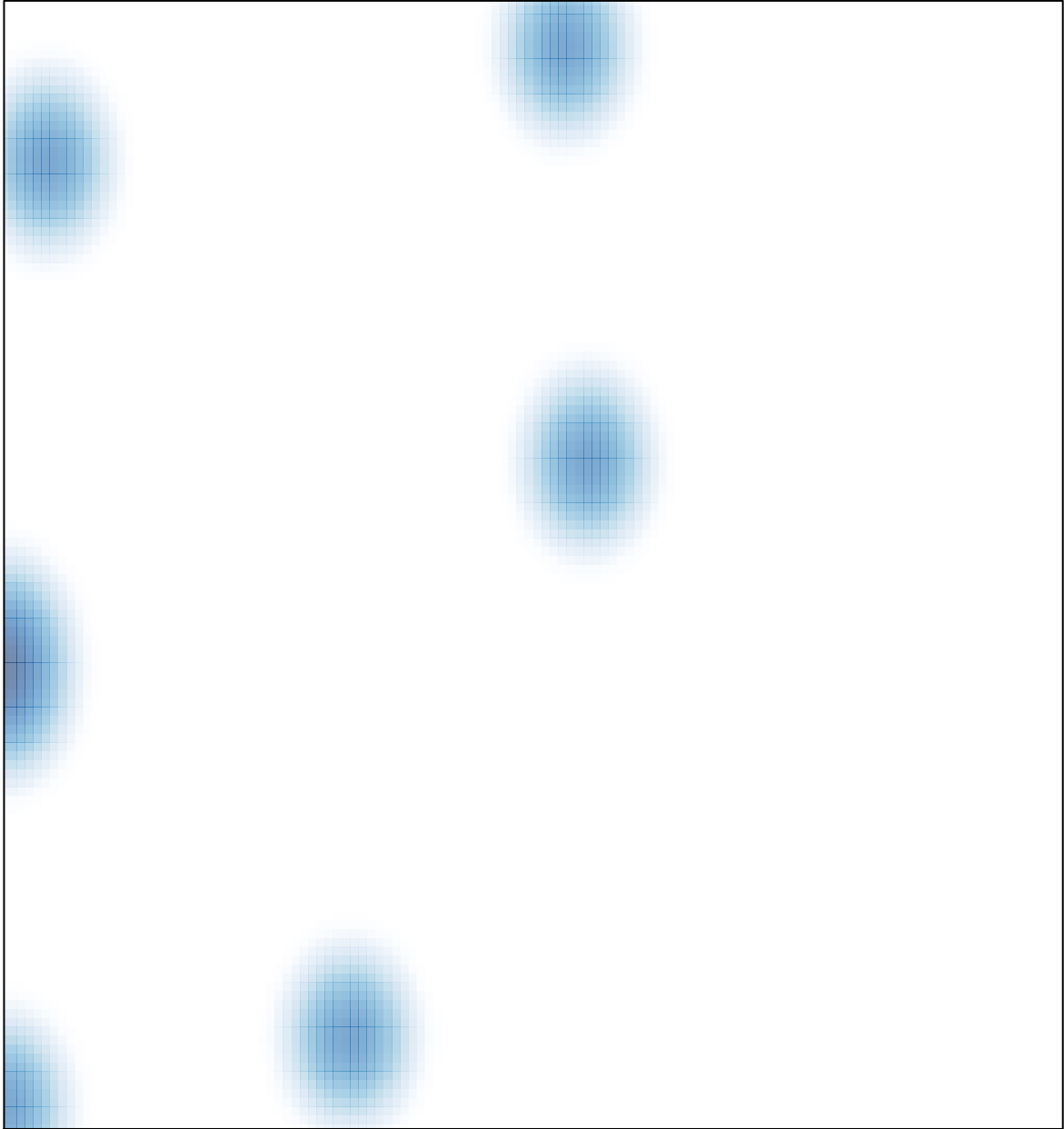


proteasome binding



features = 10
chi-square p = 0.81

proteasome binding



features = 10 , max = 1