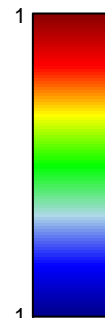
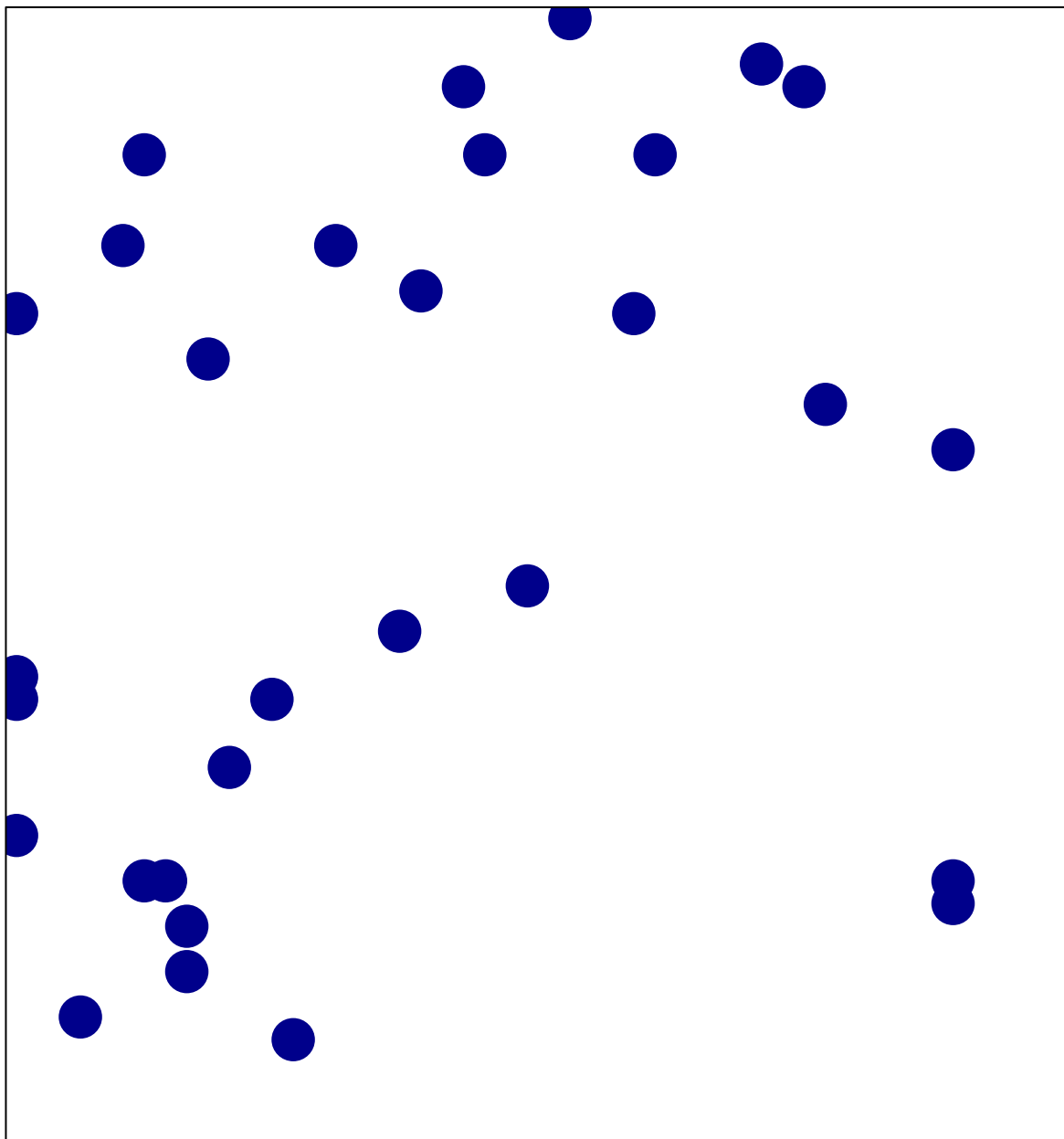


ribosome binding



features = 30
chi-square p = 0.83

ribosome binding



features = 30 , max = 1