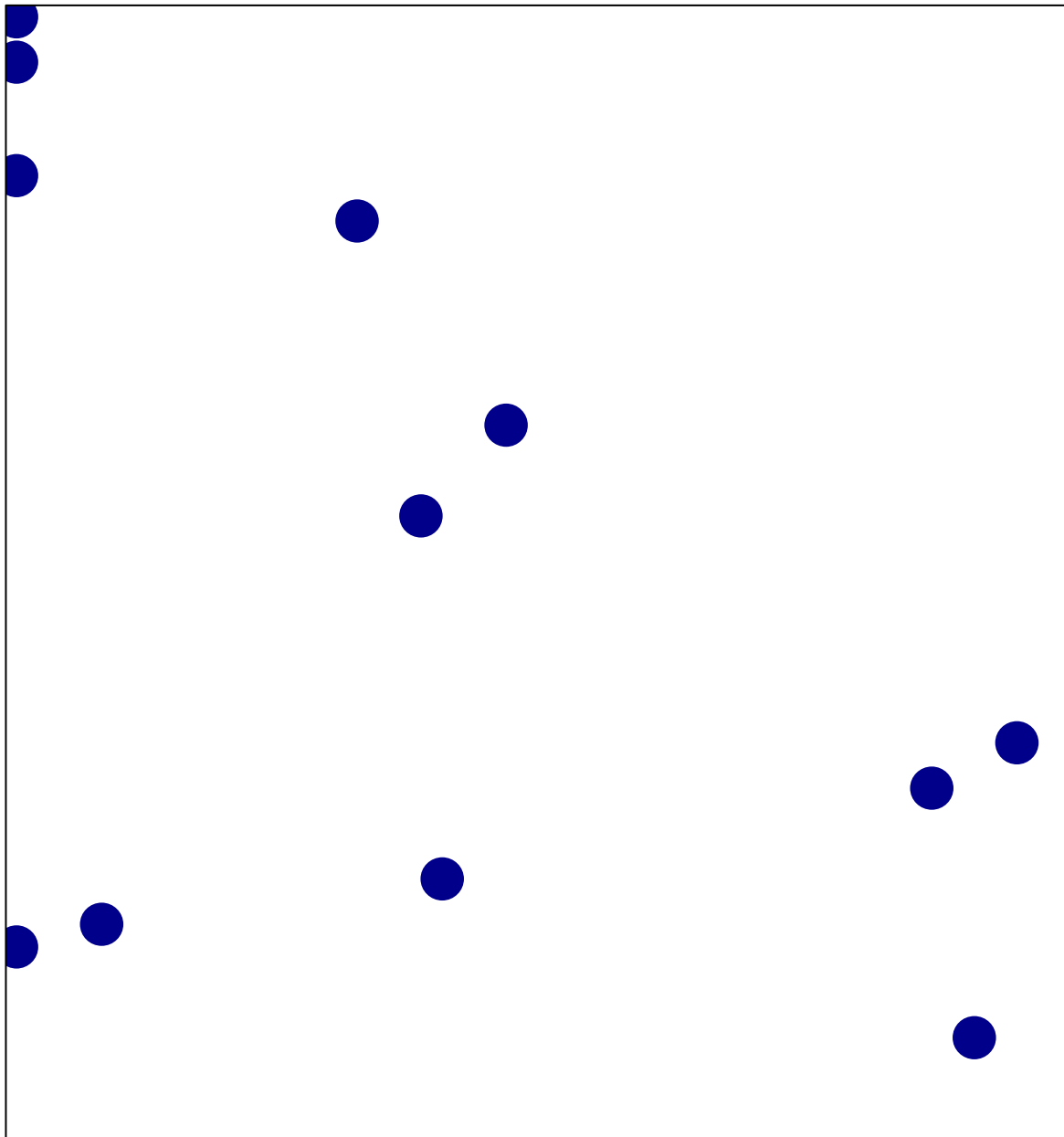
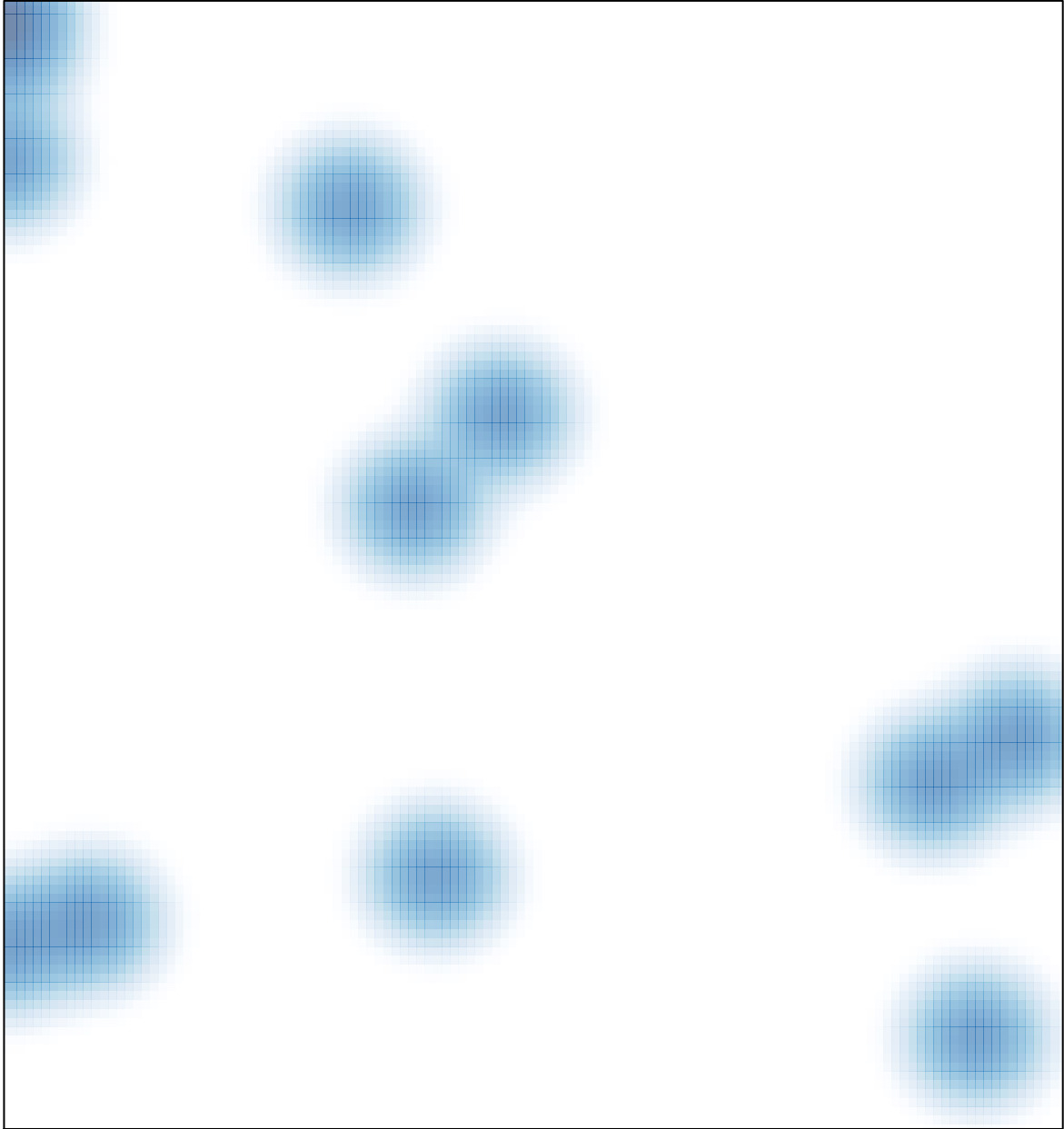


retinol binding



features = 12
chi-square p = 0.82

retinol binding



features = 12 , max = 1