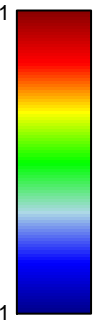
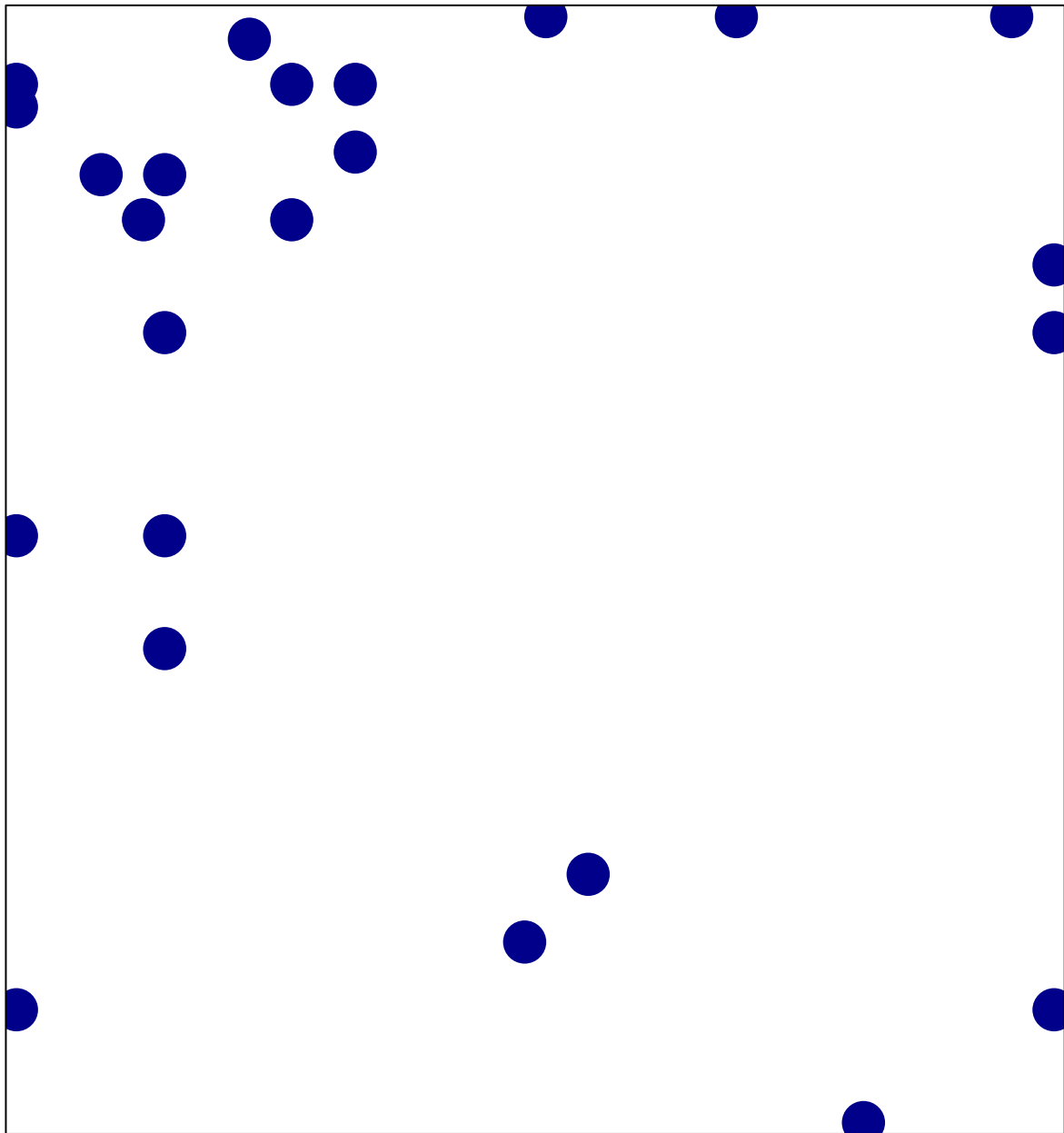


U12-type spliceosomal complex



features = 24
chi-square p = 0.8

U12-type spliceosomal complex



features = 24 , max = 1