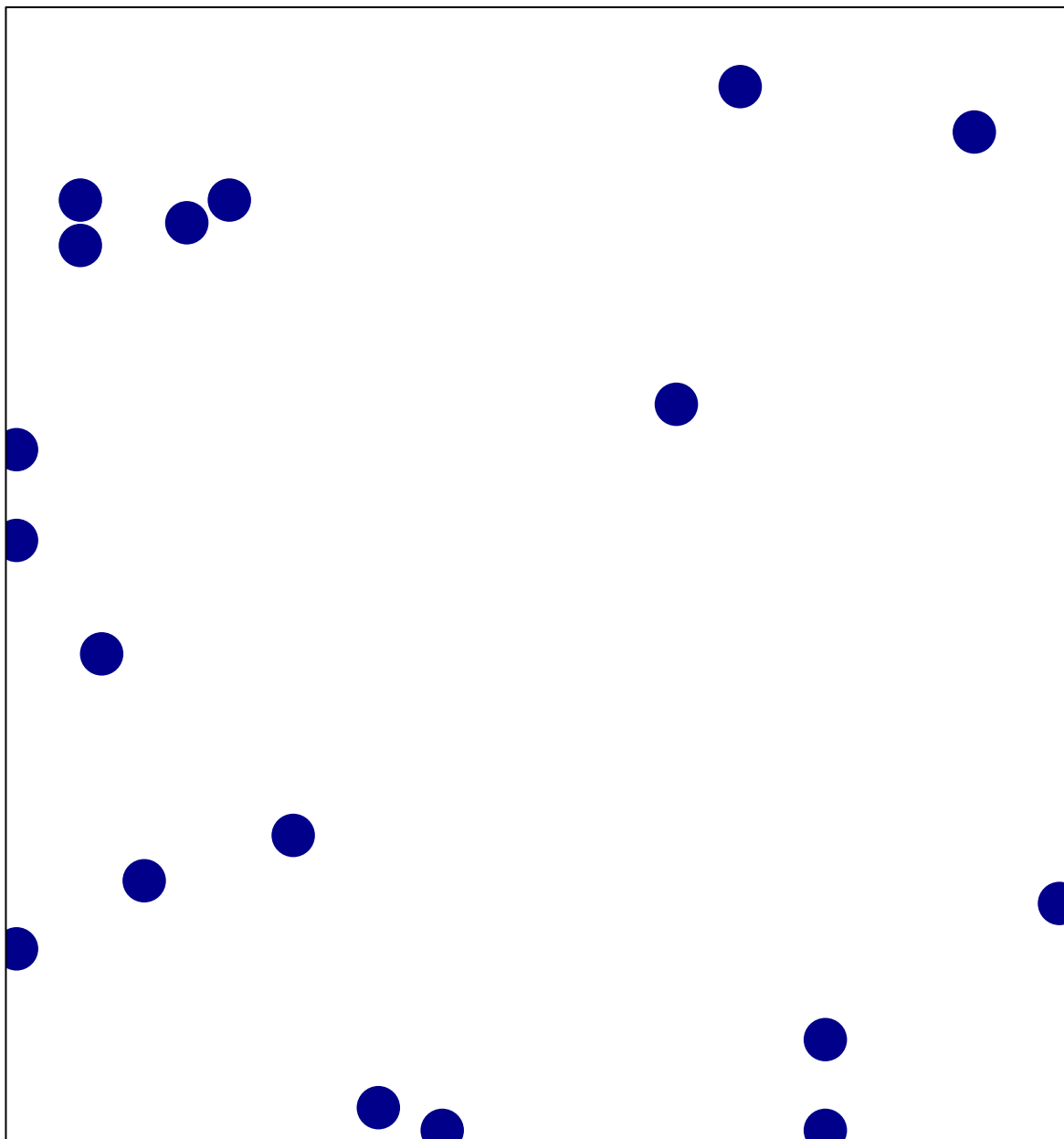
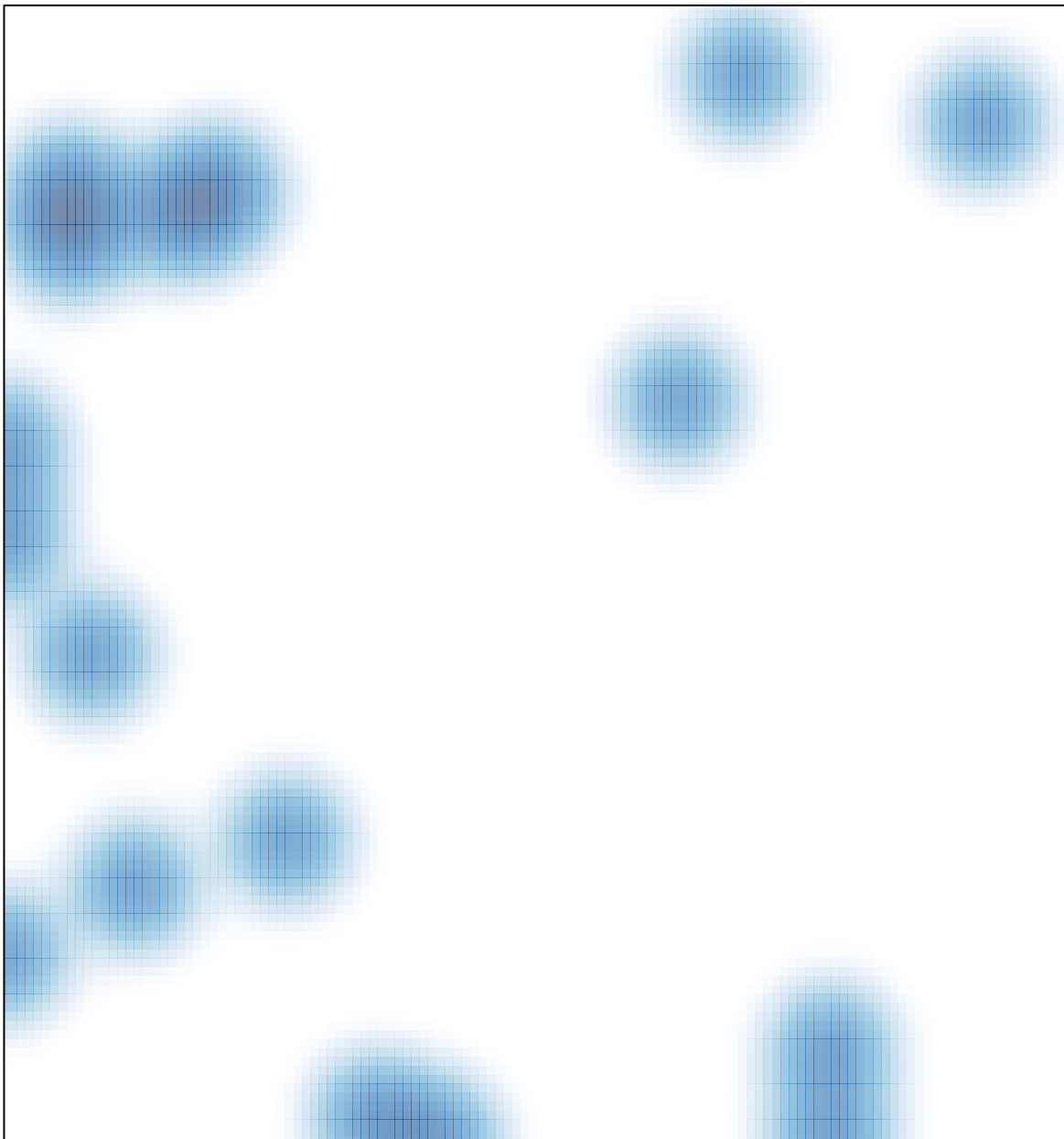


REACTOME_DEADENYLATION_OF_MRNA



features = 18
chi-square p = 0.84

REACTOME_DEADENYLATION_OF_MRNA



features = 18 , max = 1