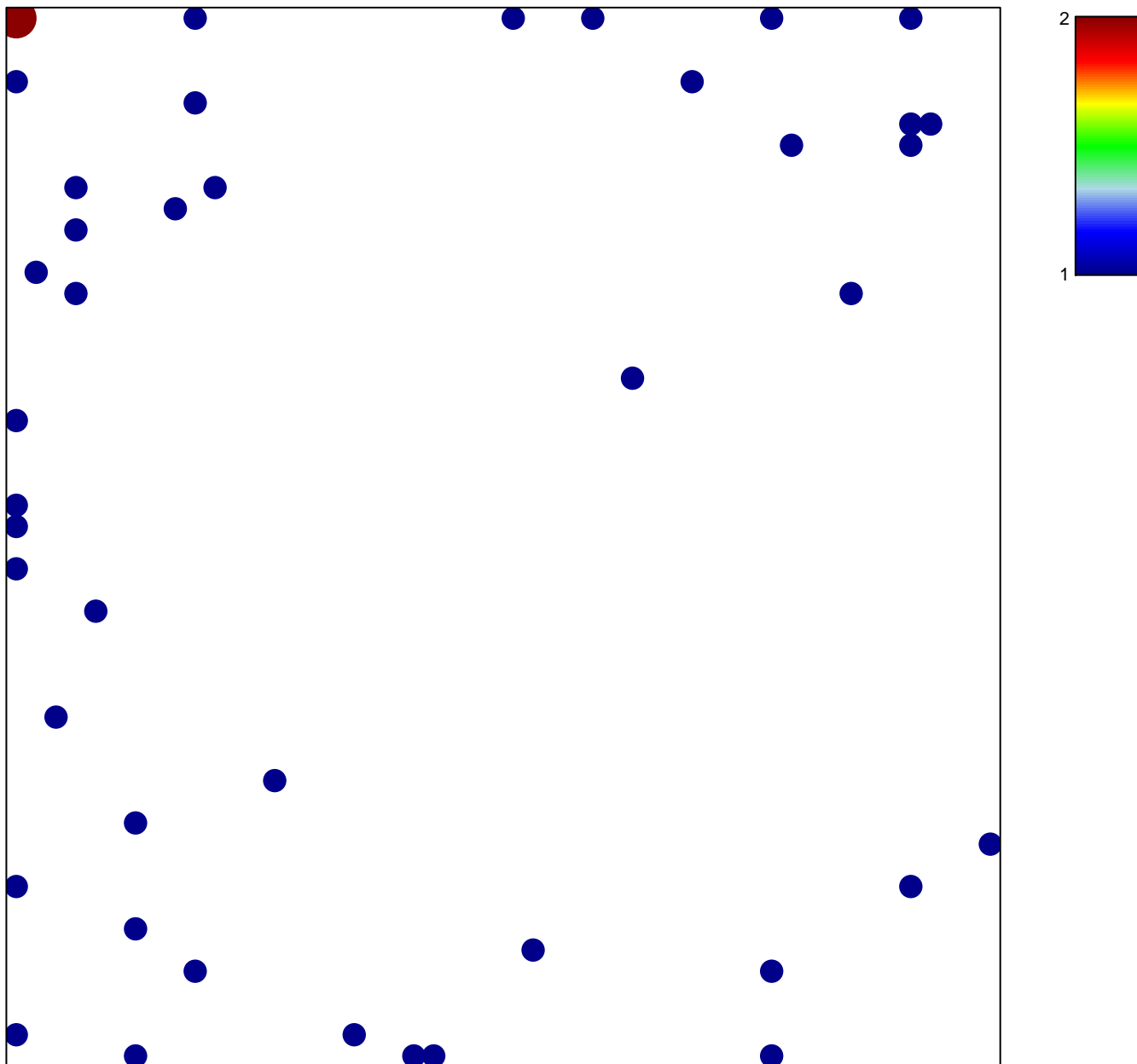
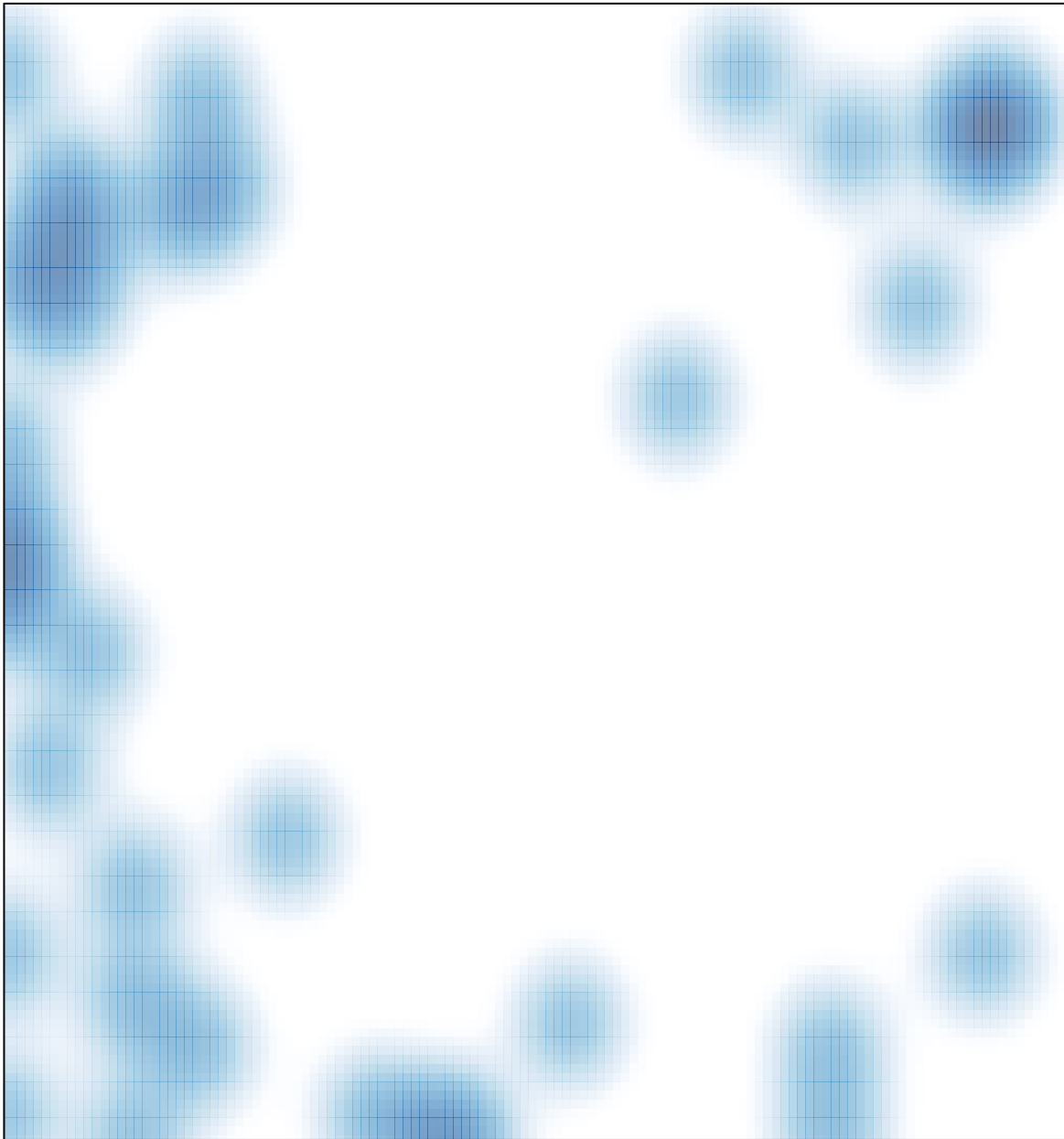


REACTOME_DEADENYLATION_DEPENDENT_MRNA_DECAY



features = 43
chi-square p = 0.77

ACTOME_DEADENYLATION_DEPENDENT_MRNA_DE



features = 43 , max = 2