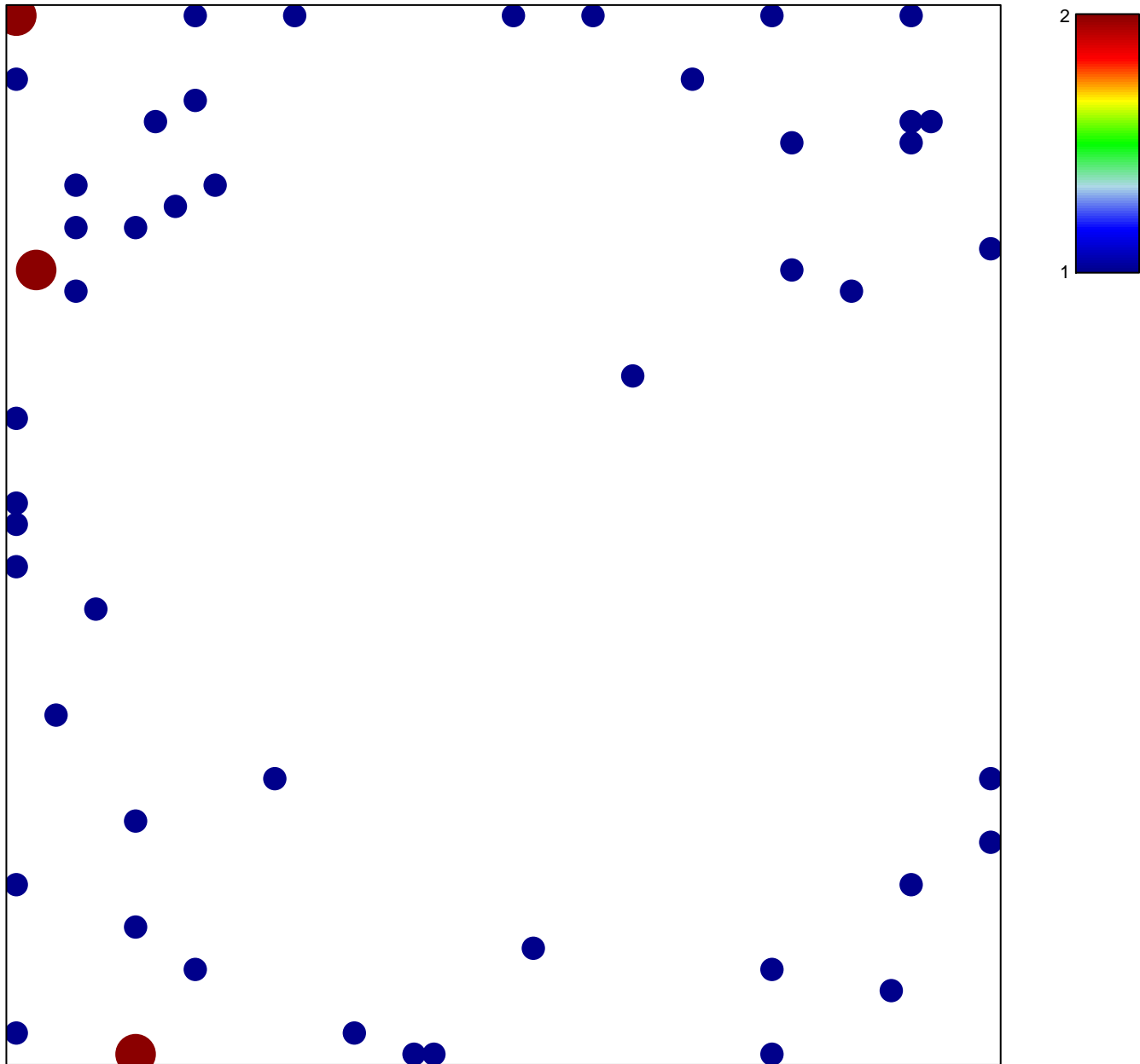
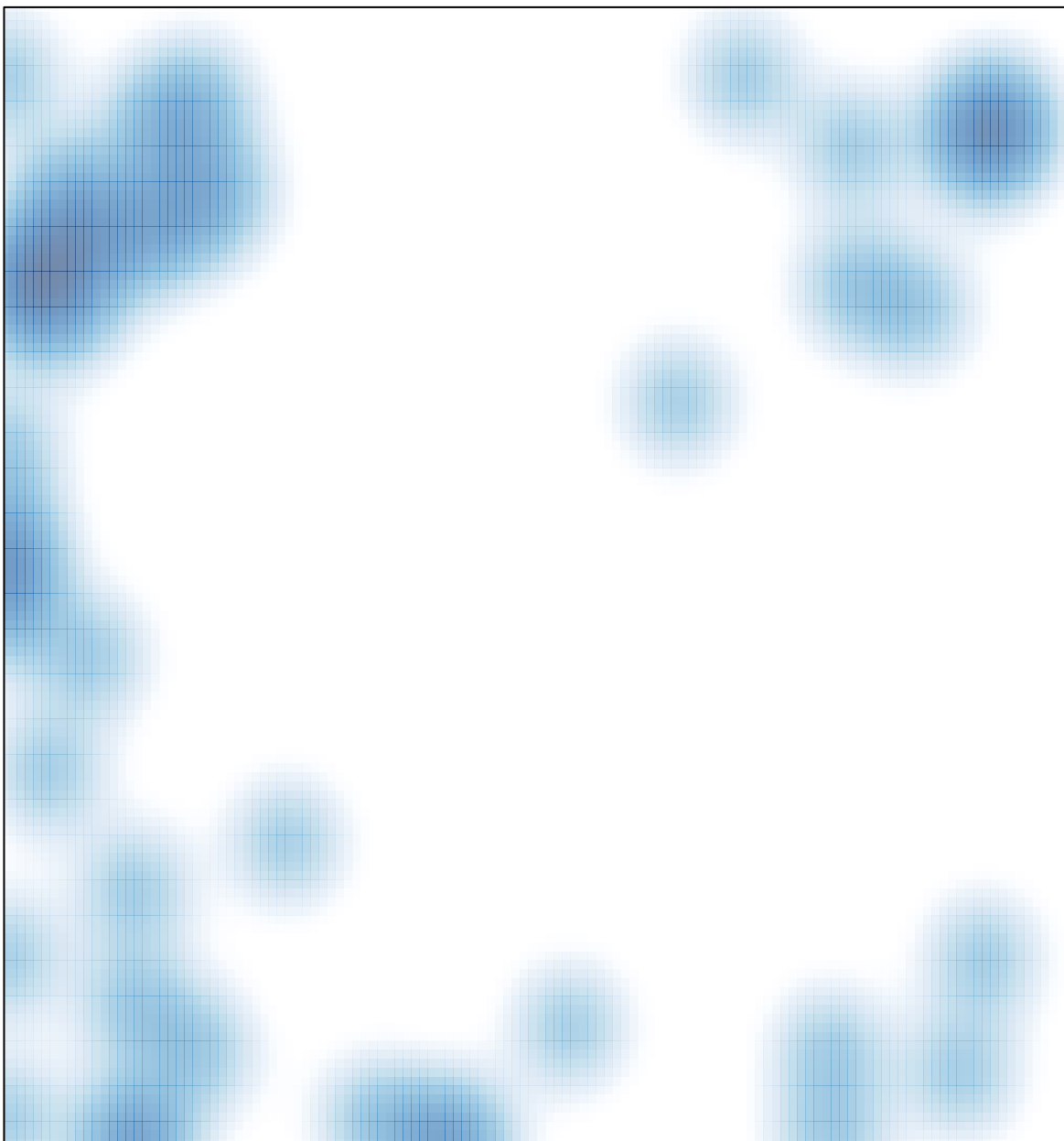


nuclear-transcribed mRNA catabolic process, deadenylation-dependent decay



features = 52
chi-square p = 0.76

scribed mRNA catabolic process, deadenylation-dep



features = 52 , max = 2