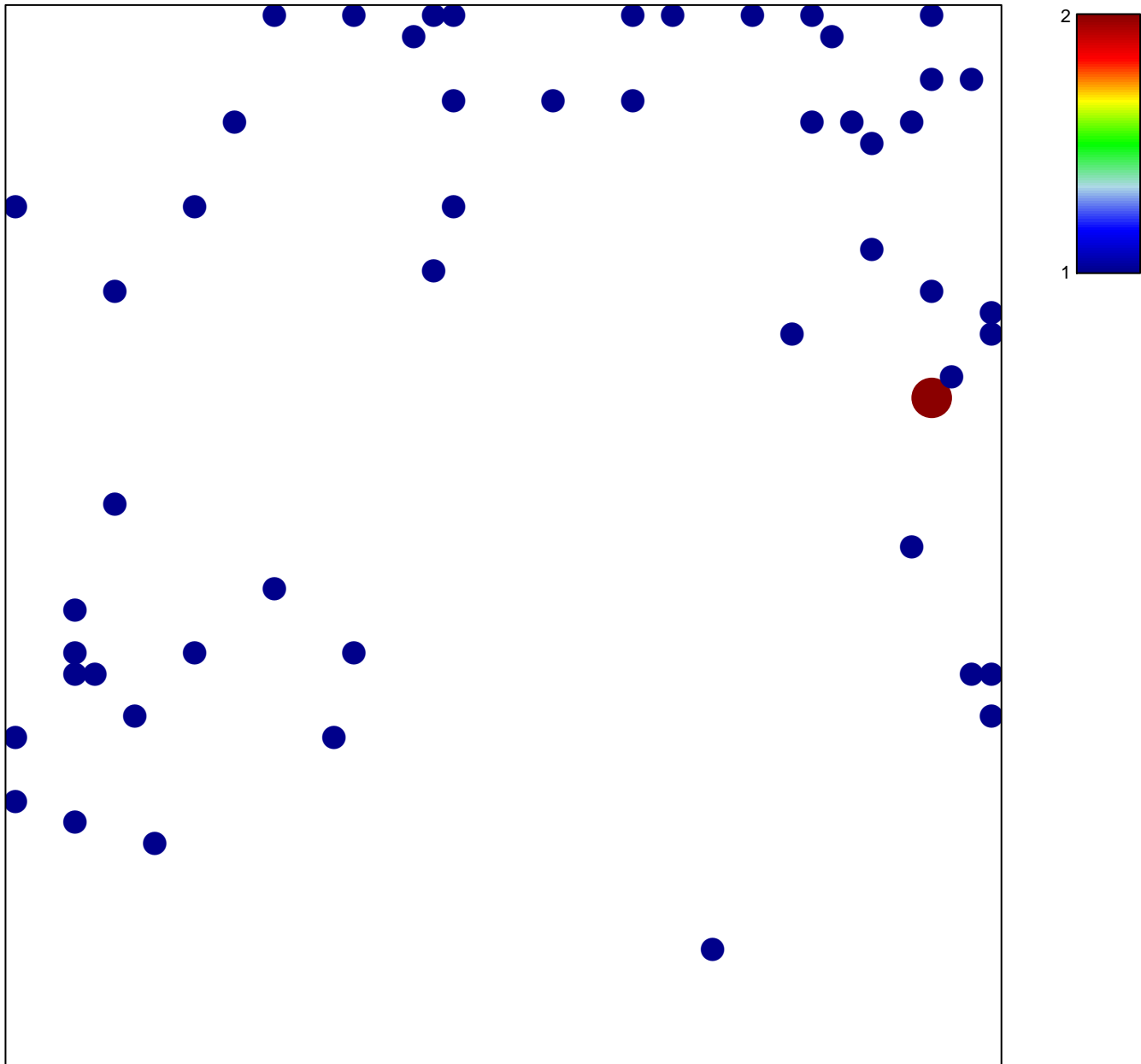
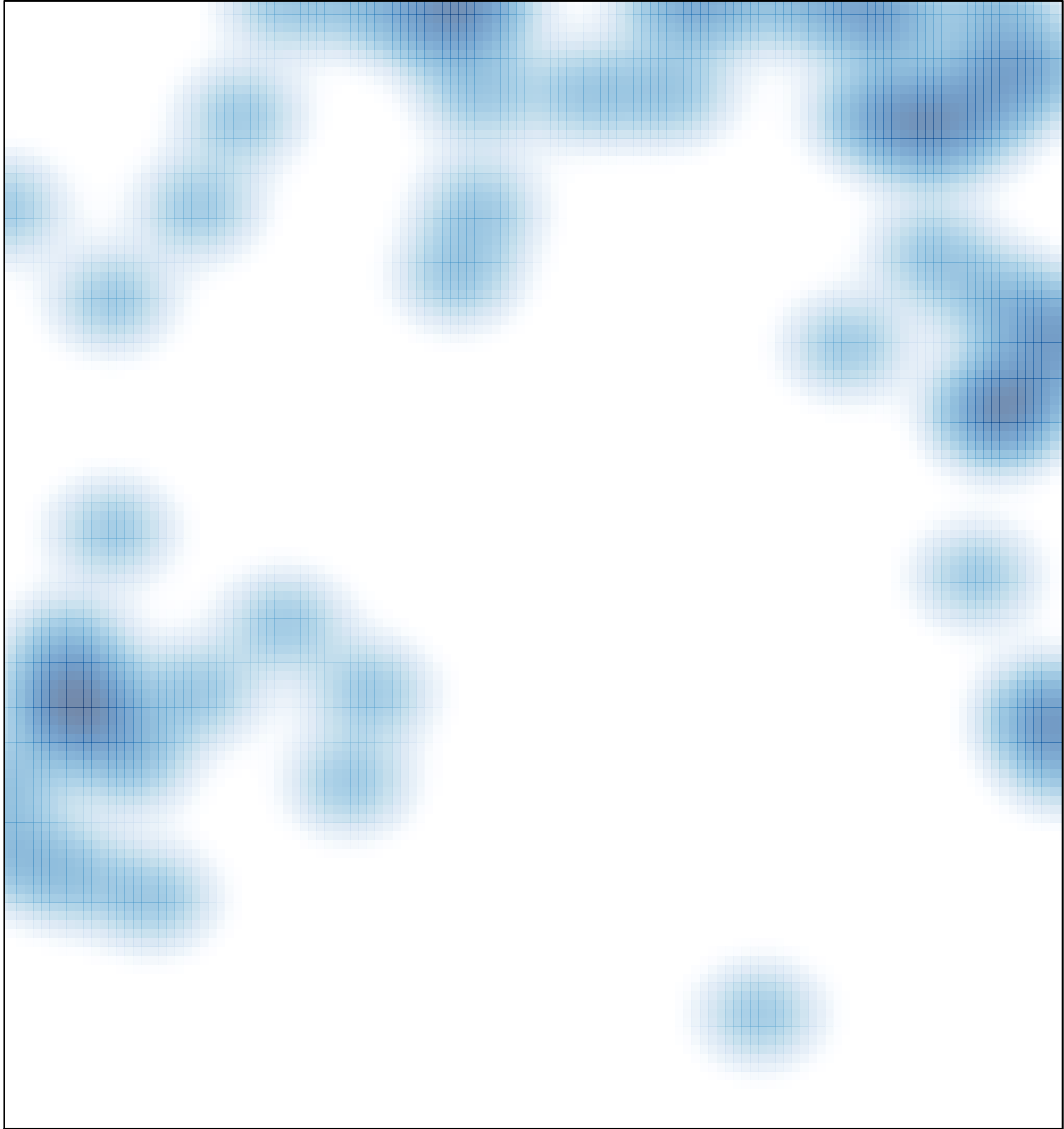


nuclear-transcribed mRNA catabolic process, deadenylation-dependent decay



features = 53
chi-square p = 0.78

scribed mRNA catabolic process, deadenylation-dep



features = 53 , max = 2