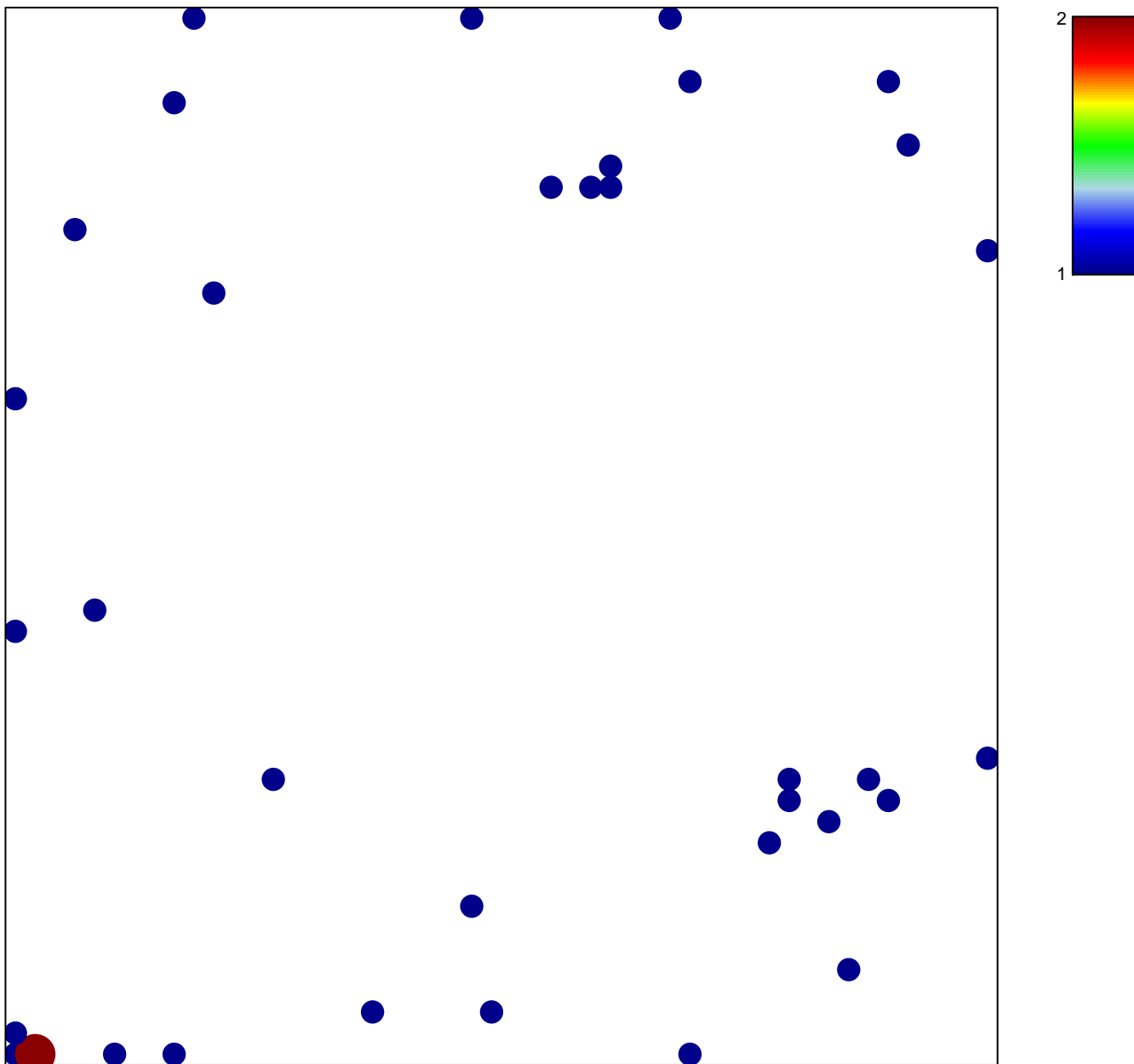
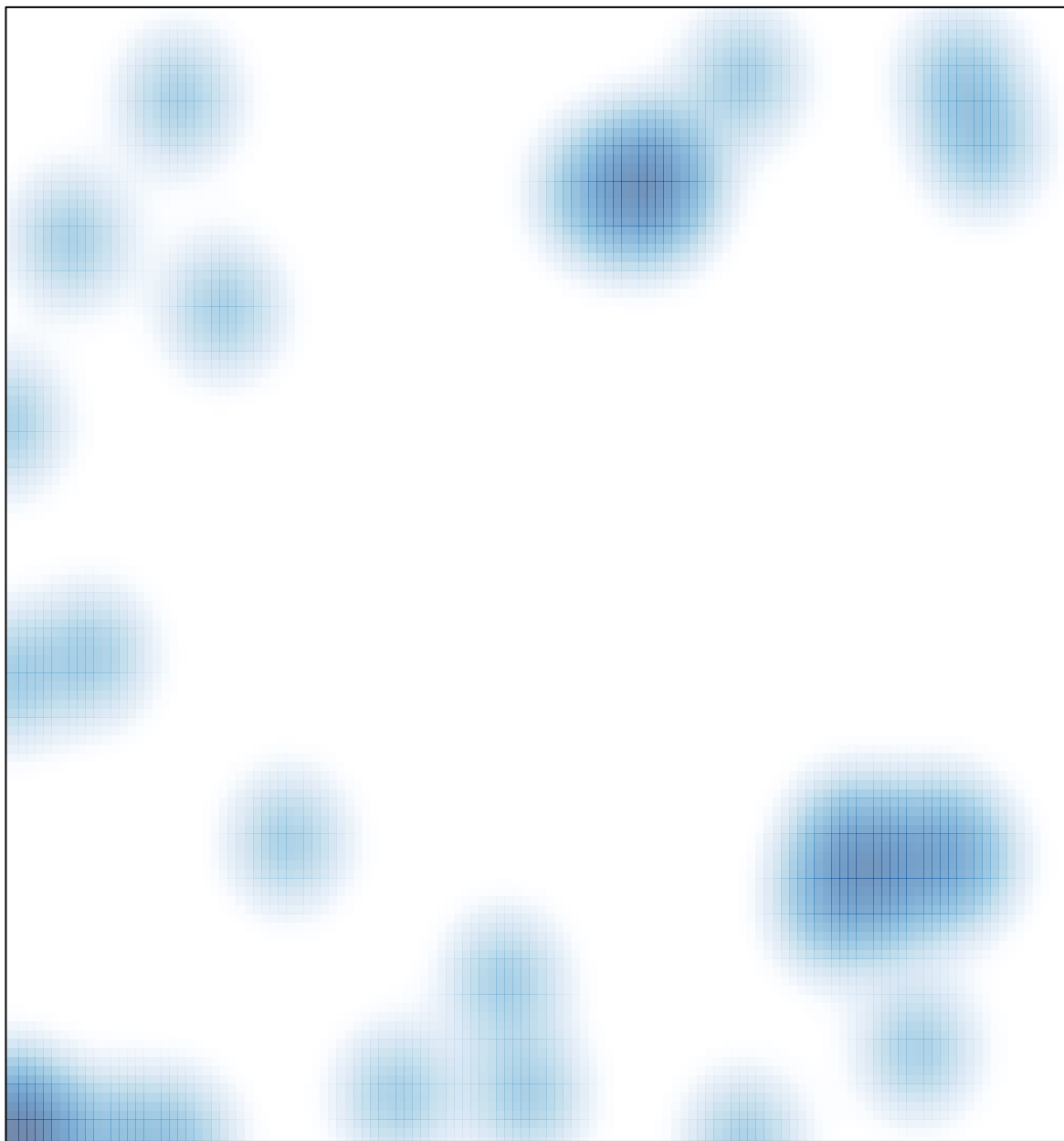


nuclear-transcribed mRNA catabolic process, nonsense-mediated decay



features = 36
chi-square p = 0.81

transcribed mRNA catabolic process, nonsense-med



features = 36 , max = 2