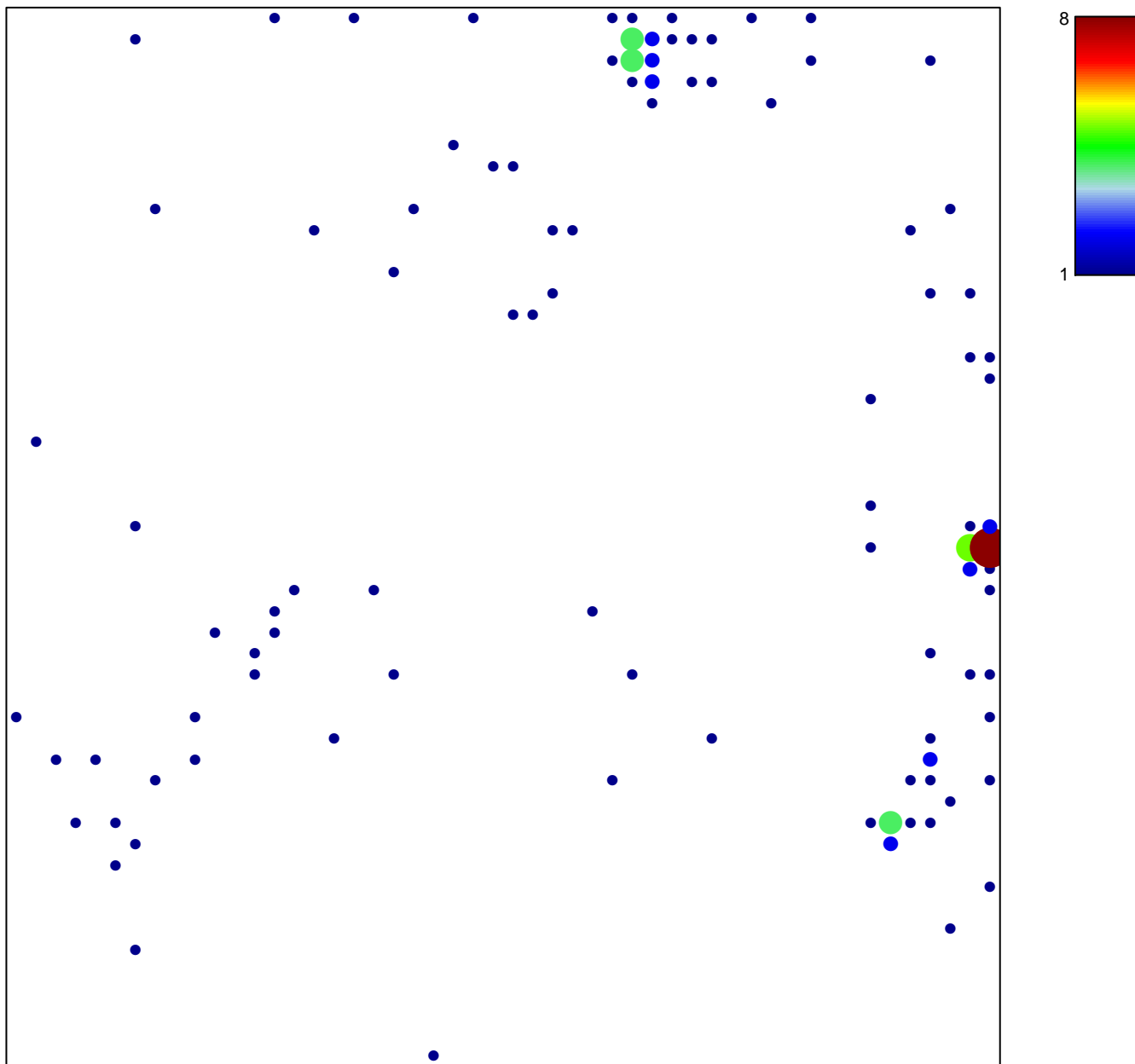
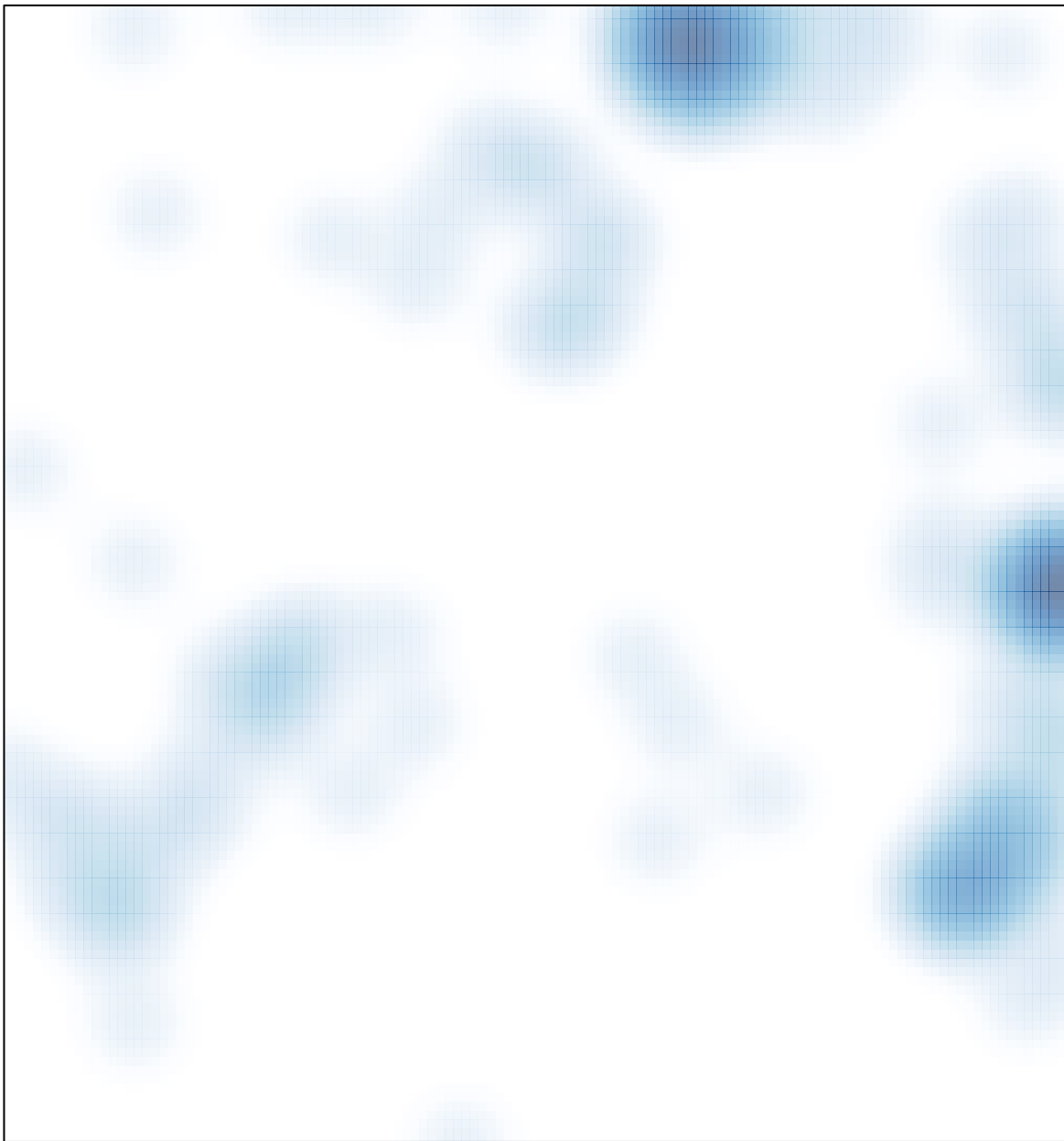


nuclear-transcribed mRNA catabolic process, nonsense-mediated decay



features = 125
chi-square p = 0.69

transcribed mRNA catabolic process, nonsense-med



features = 125 , max = 8