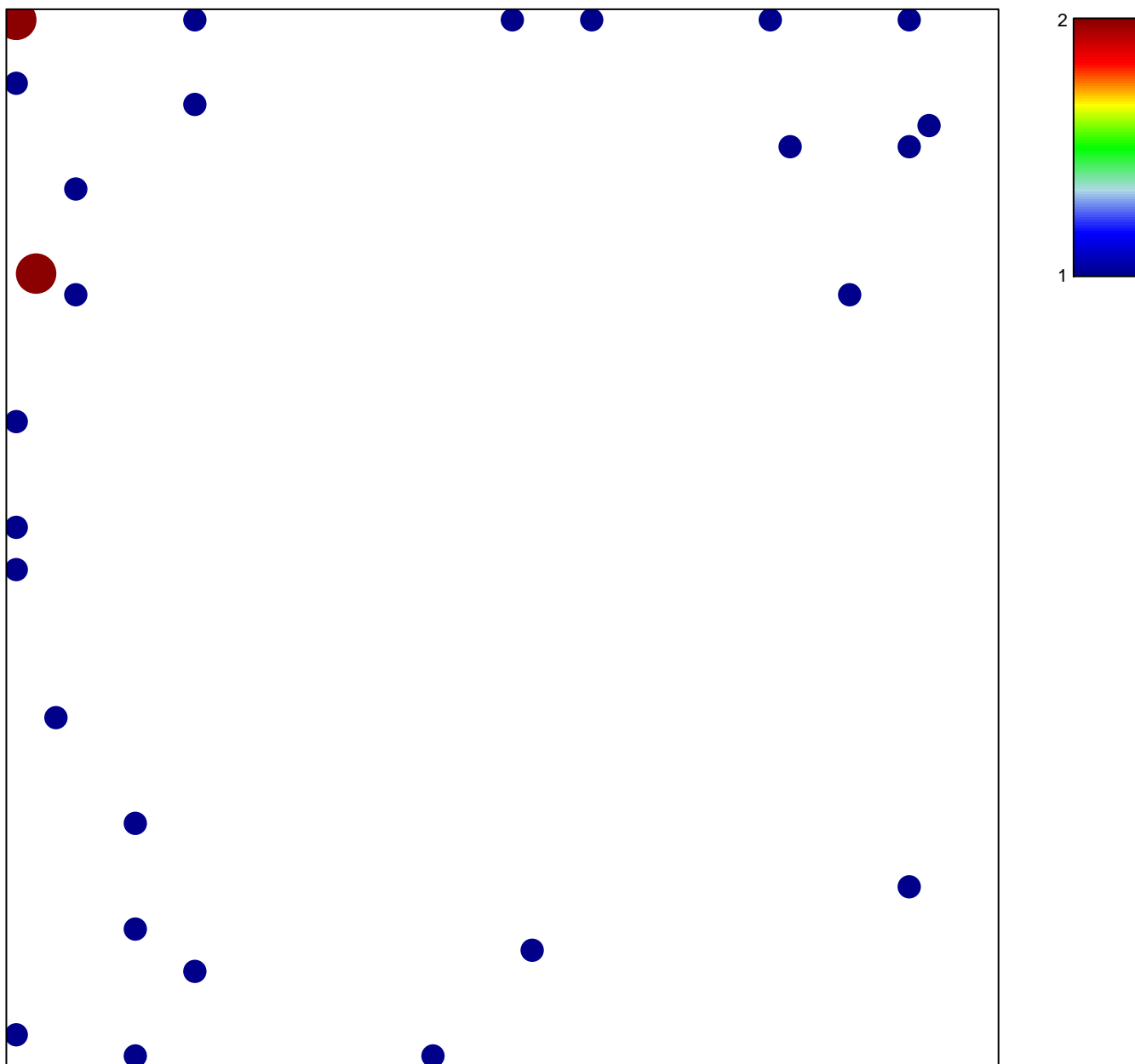
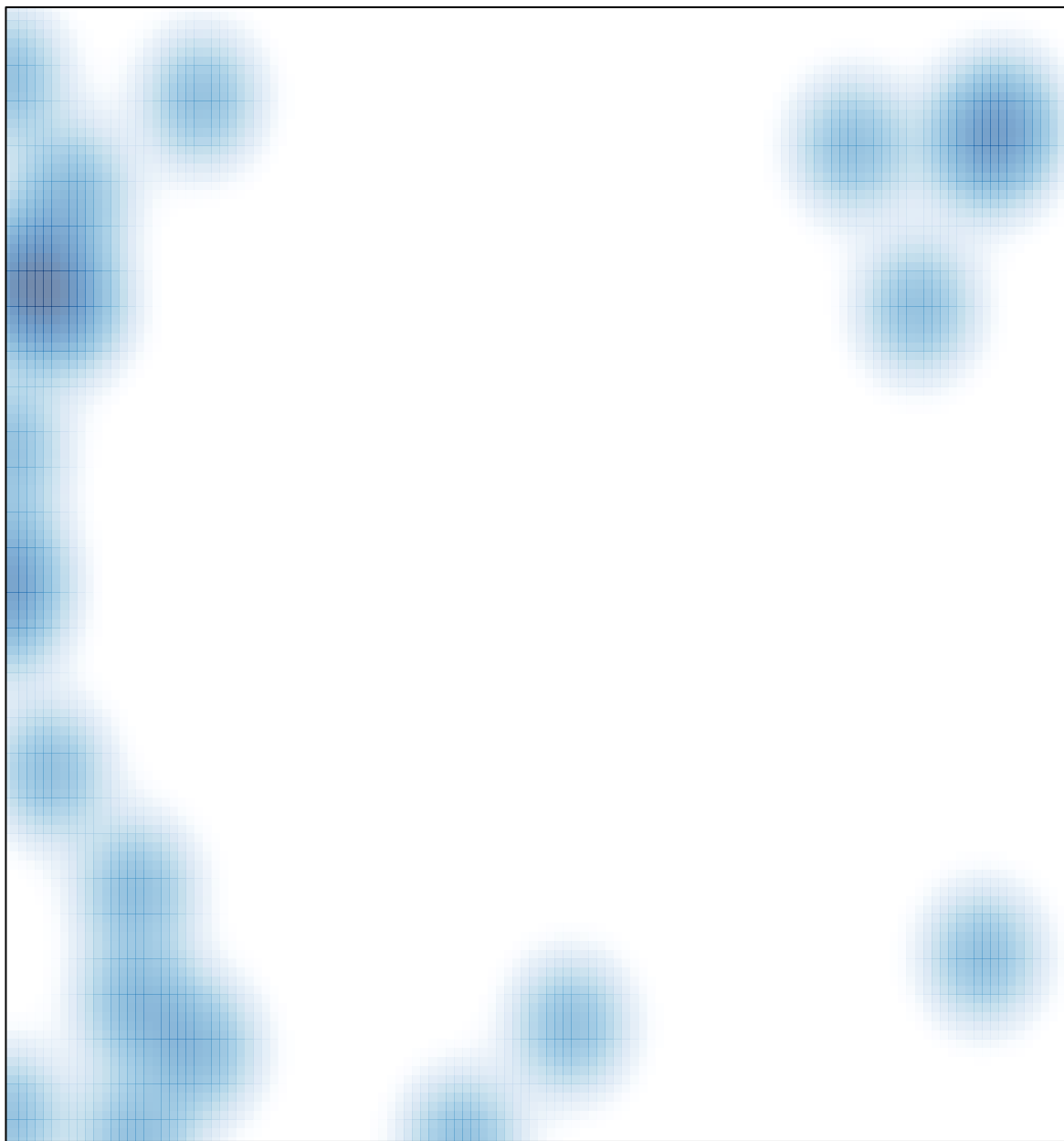


cleolytic nuclear-transcribed mRNA catabolic process involved in deadenylation-depende



features = 29
chi-square p = 0.76

transcribed mRNA catabolic process involved in dead



features = 29 , max = 2