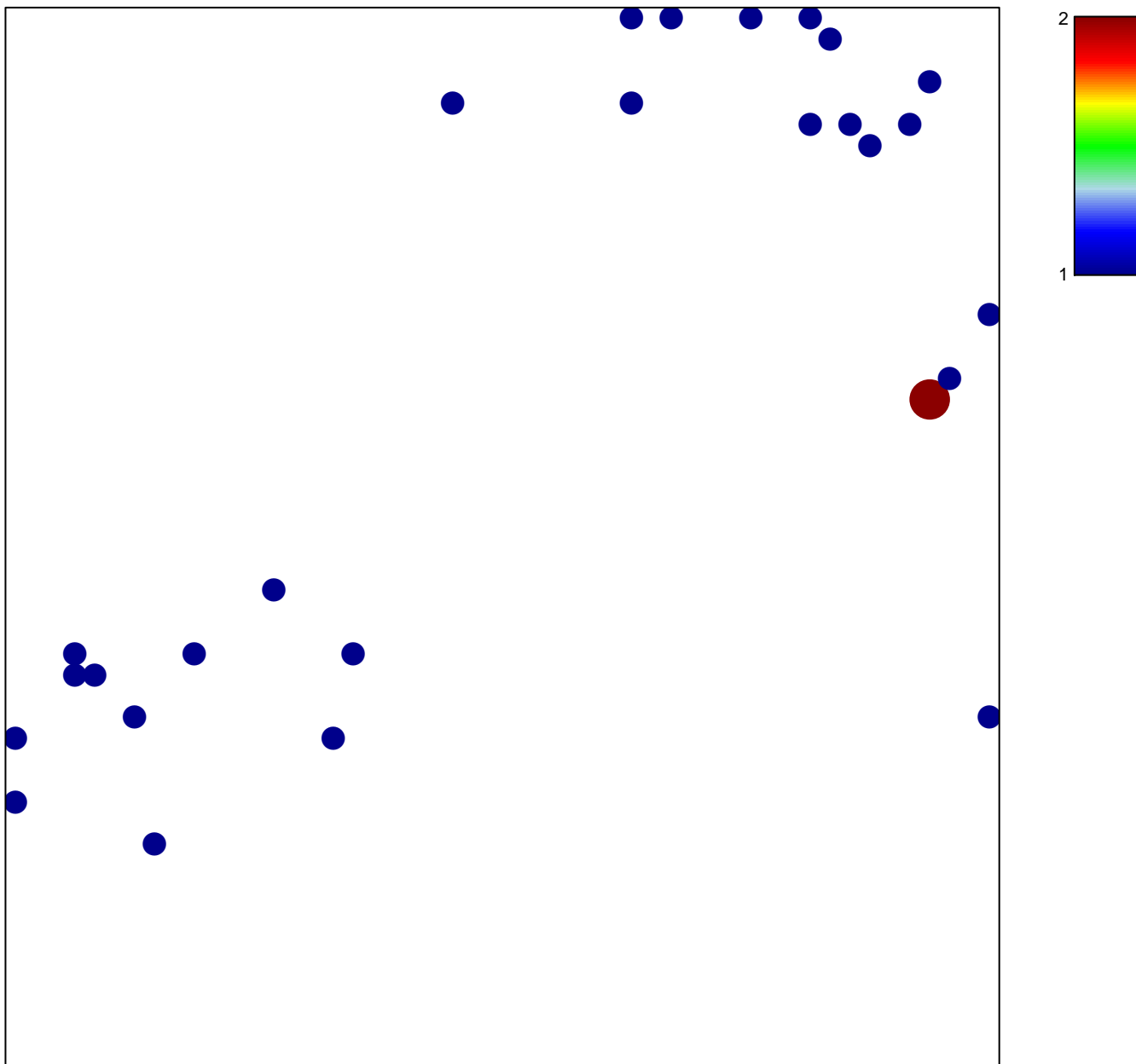
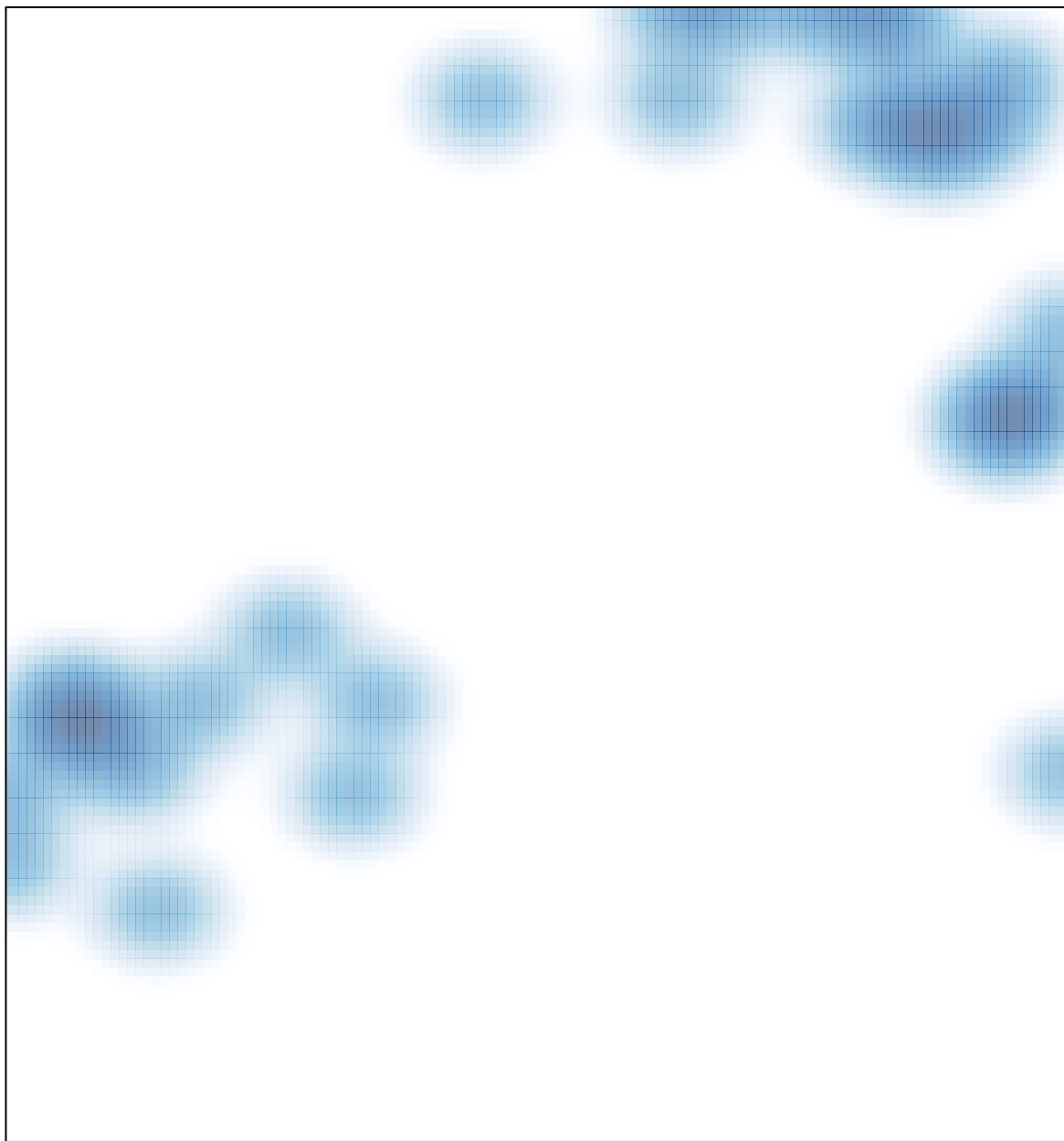


cleolytic nuclear-transcribed mRNA catabolic process involved in deadenylation-depende



features = 28
chi-square p = 0.82

transcribed mRNA catabolic process involved in dead



features = 28 , max = 2