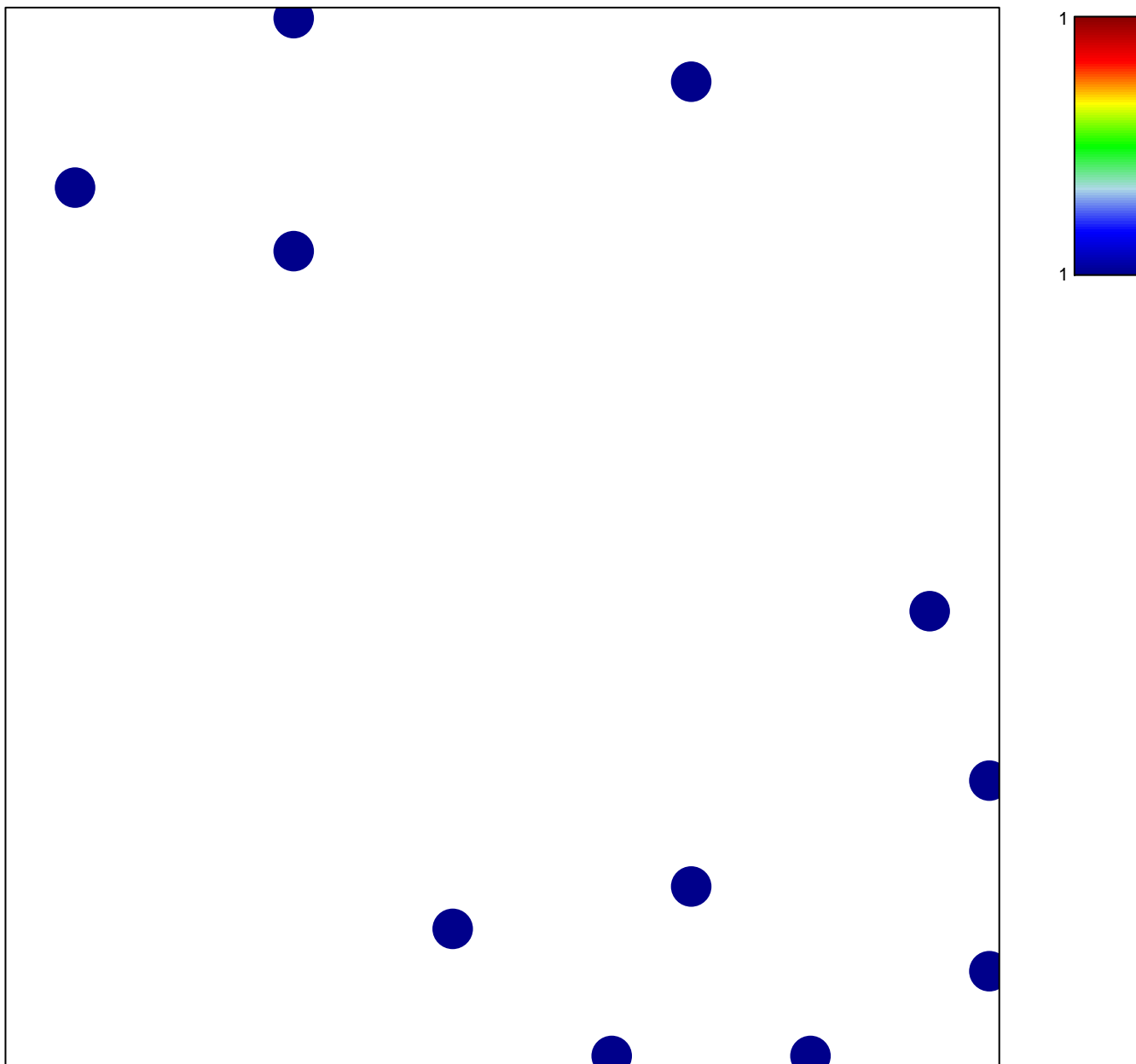
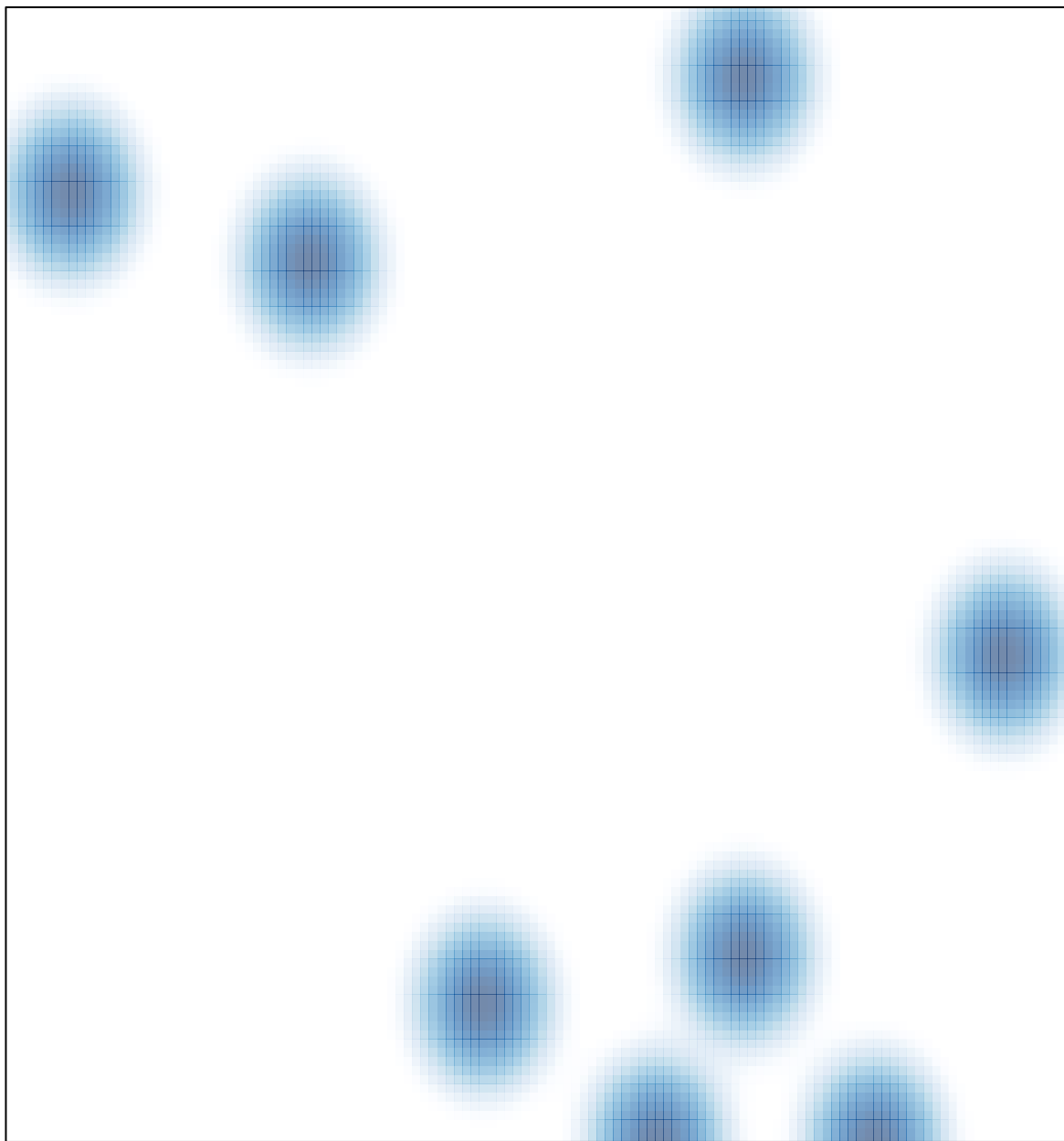


positive regulation of nuclear-transcribed mRNA catabolic process, deadenylation-dependent



features = 11
chi-square p = 0.83

nuclear-transcribed mRNA catabolic process, deaden



features = 11 , max = 1