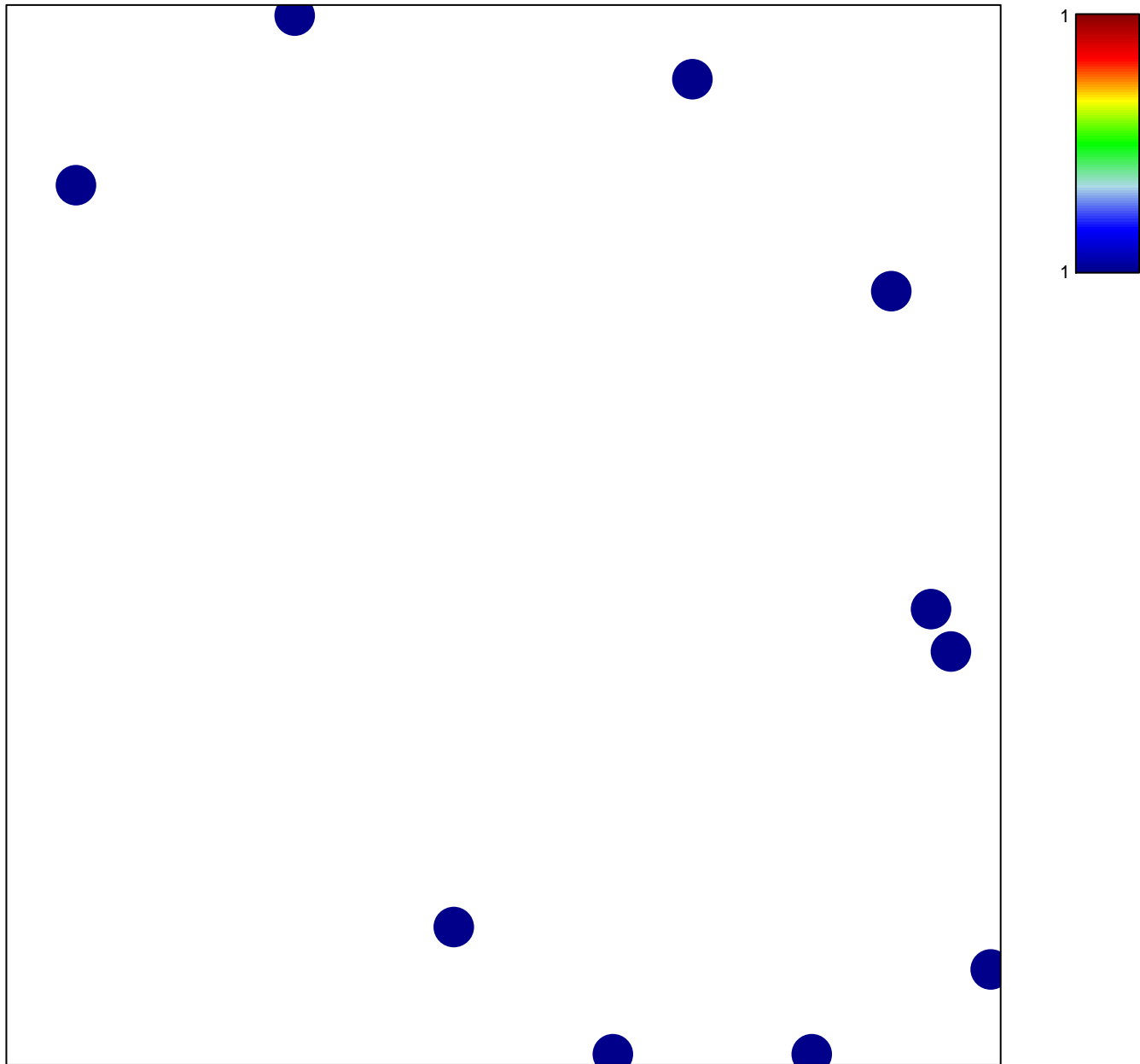
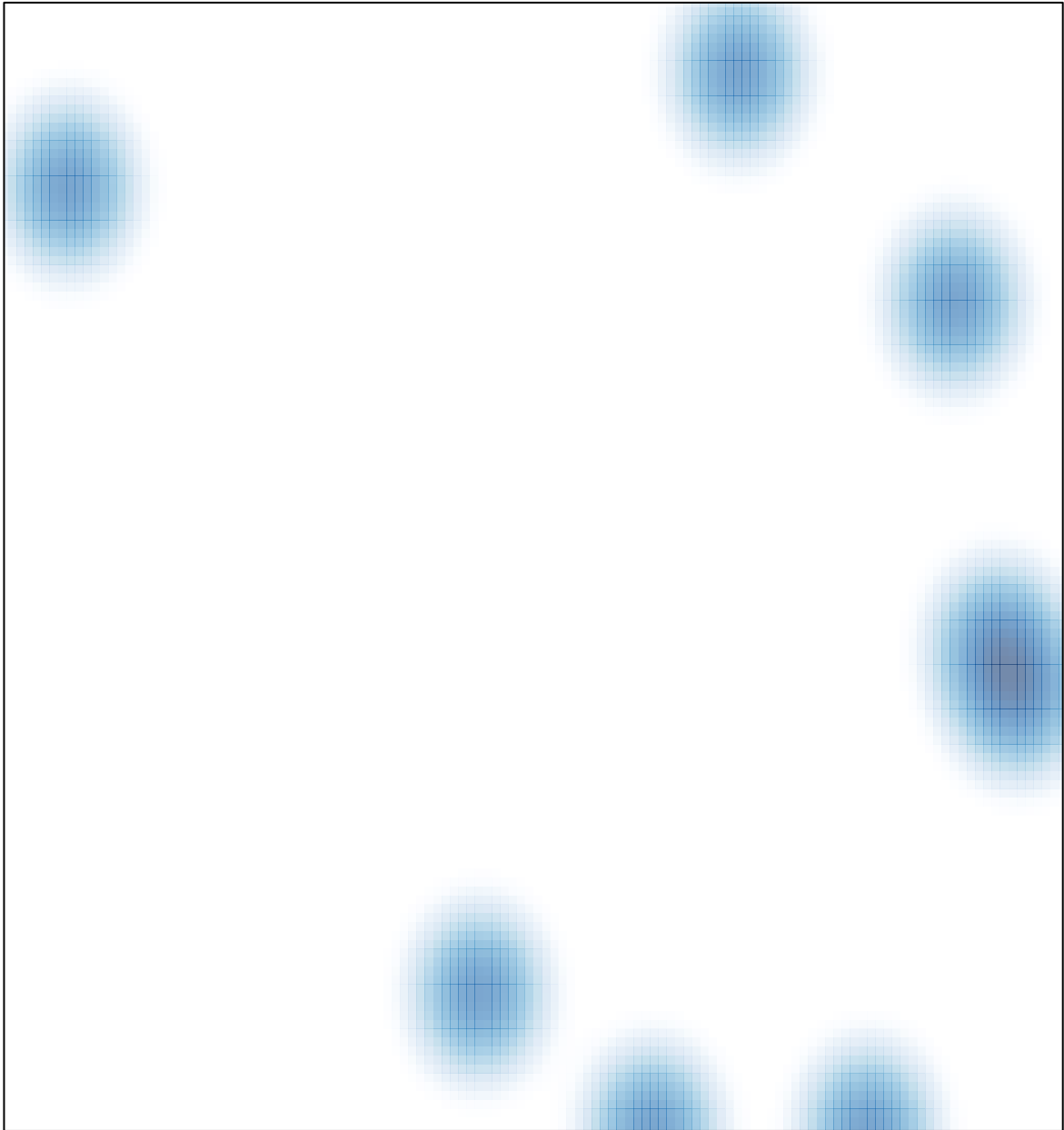


positive regulation of nuclear-transcribed mRNA poly(A) tail shortening



features = 10
chi-square p = 0.85

regulation of nuclear-transcribed mRNA poly(A) tail s



features = 10 , max = 1