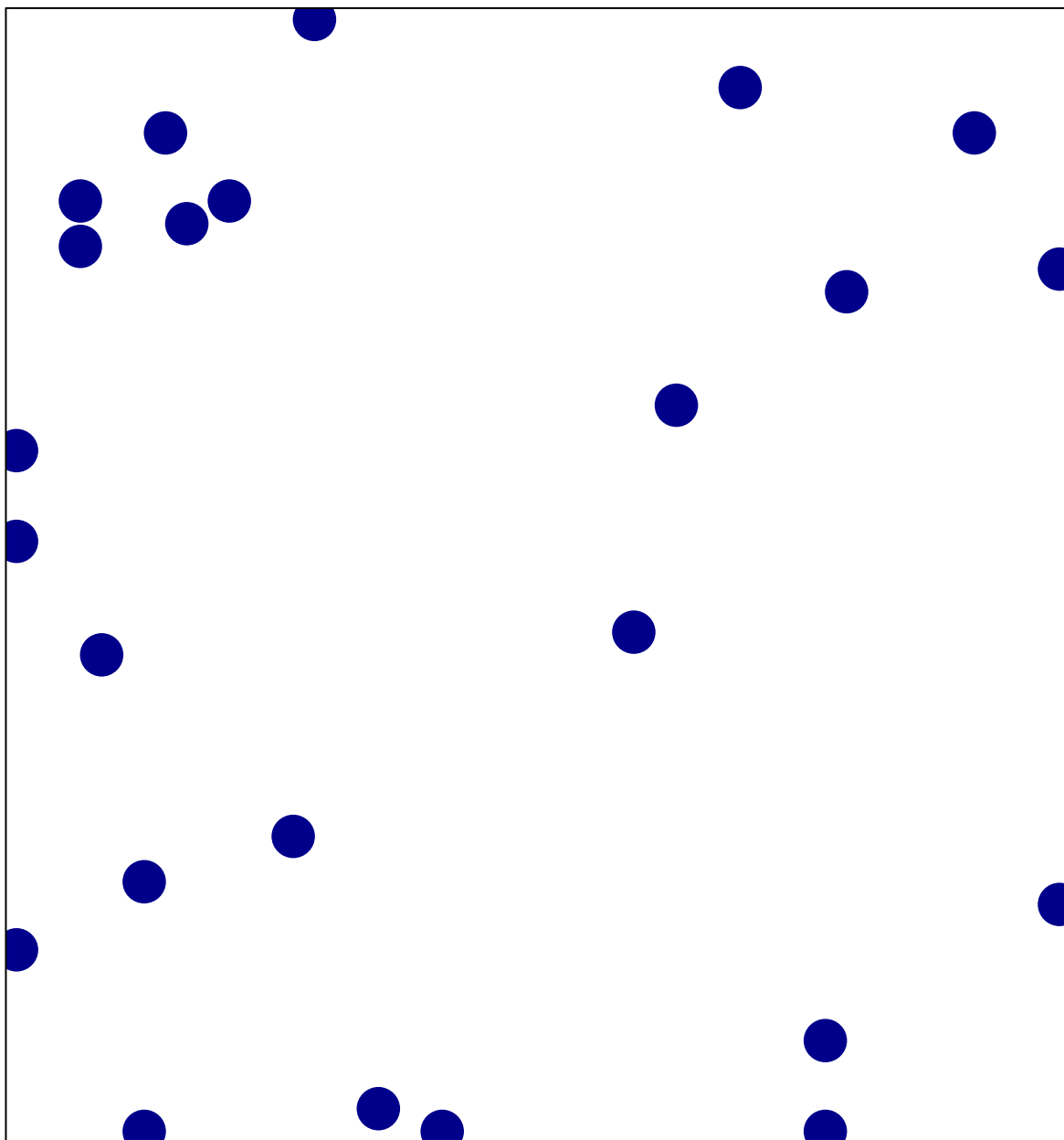
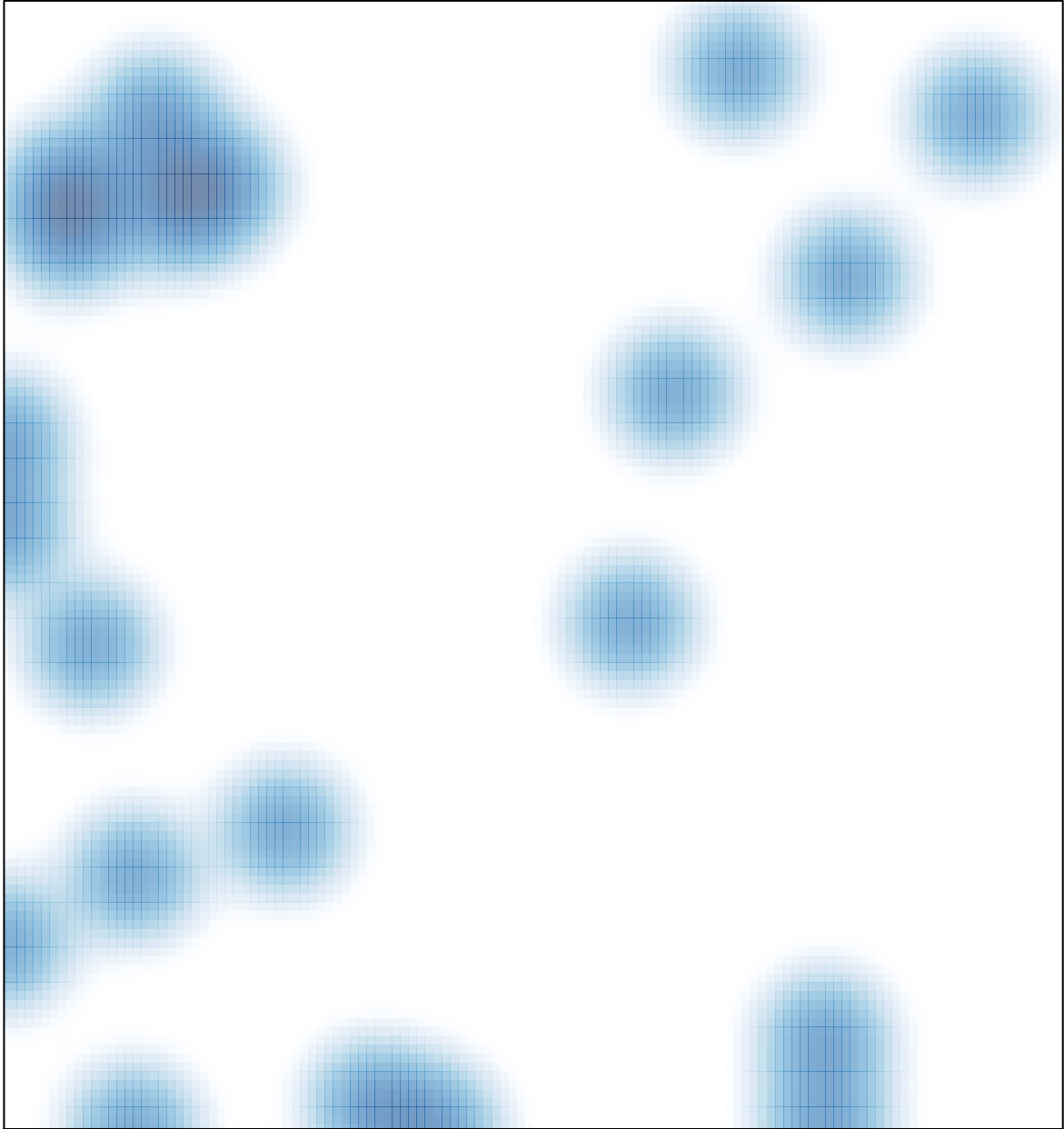


# nuclear-transcribed mRNA poly(A) tail shortening



# features = 24  
chi-square p = 0.84

# nuclear-transcribed mRNA poly(A) tail shortening



# features = 24 , max = 1