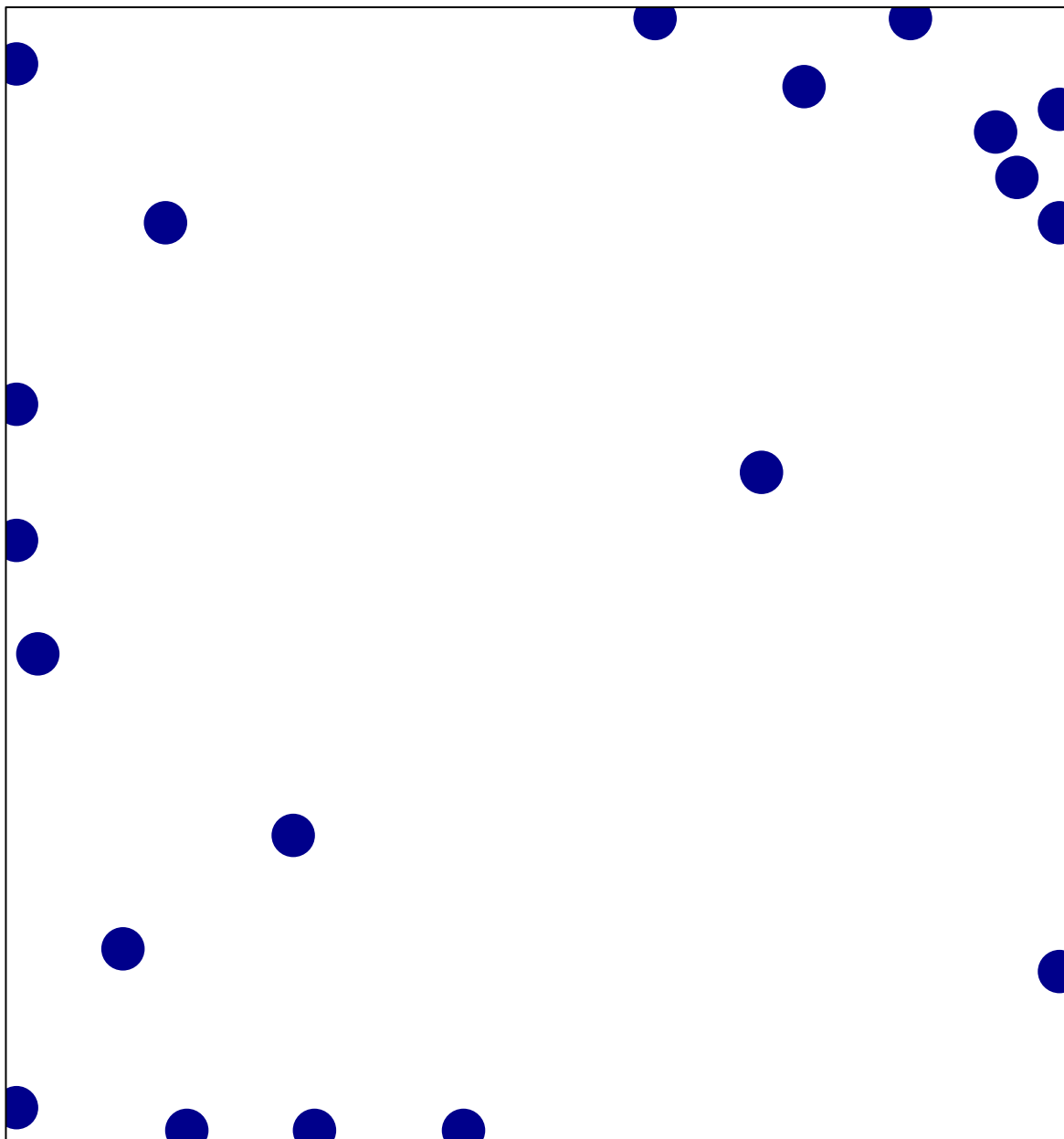
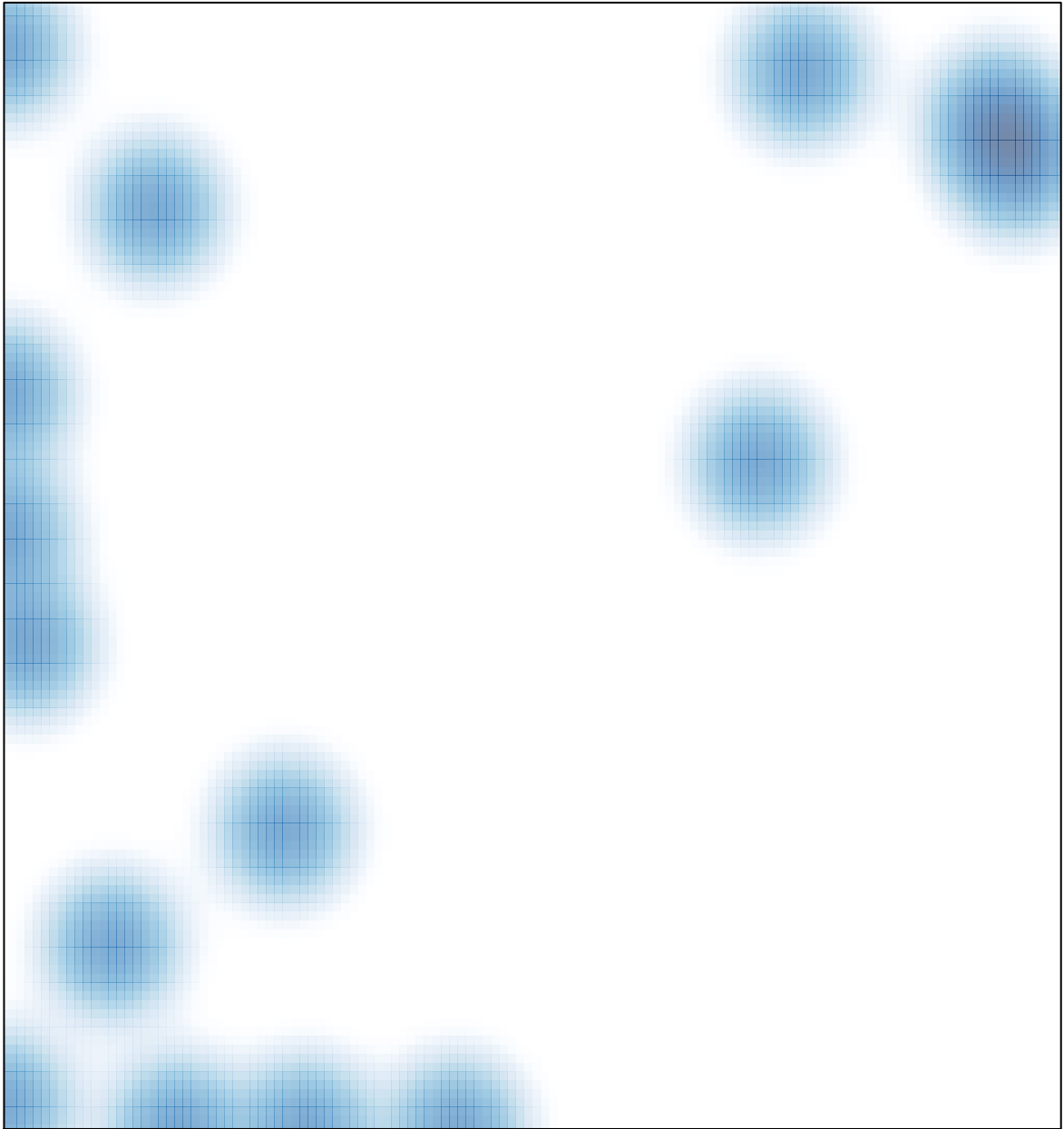


# REACTOME\_RNA\_POL\_I\_TRANSCRIPTION\_TERMINATION



# features = 20  
chi-square p = 0.82

# FACTOME\_RNA\_POL\_I\_TRANSCRIPTION\_TERMINAT



# features = 20 , max = 1