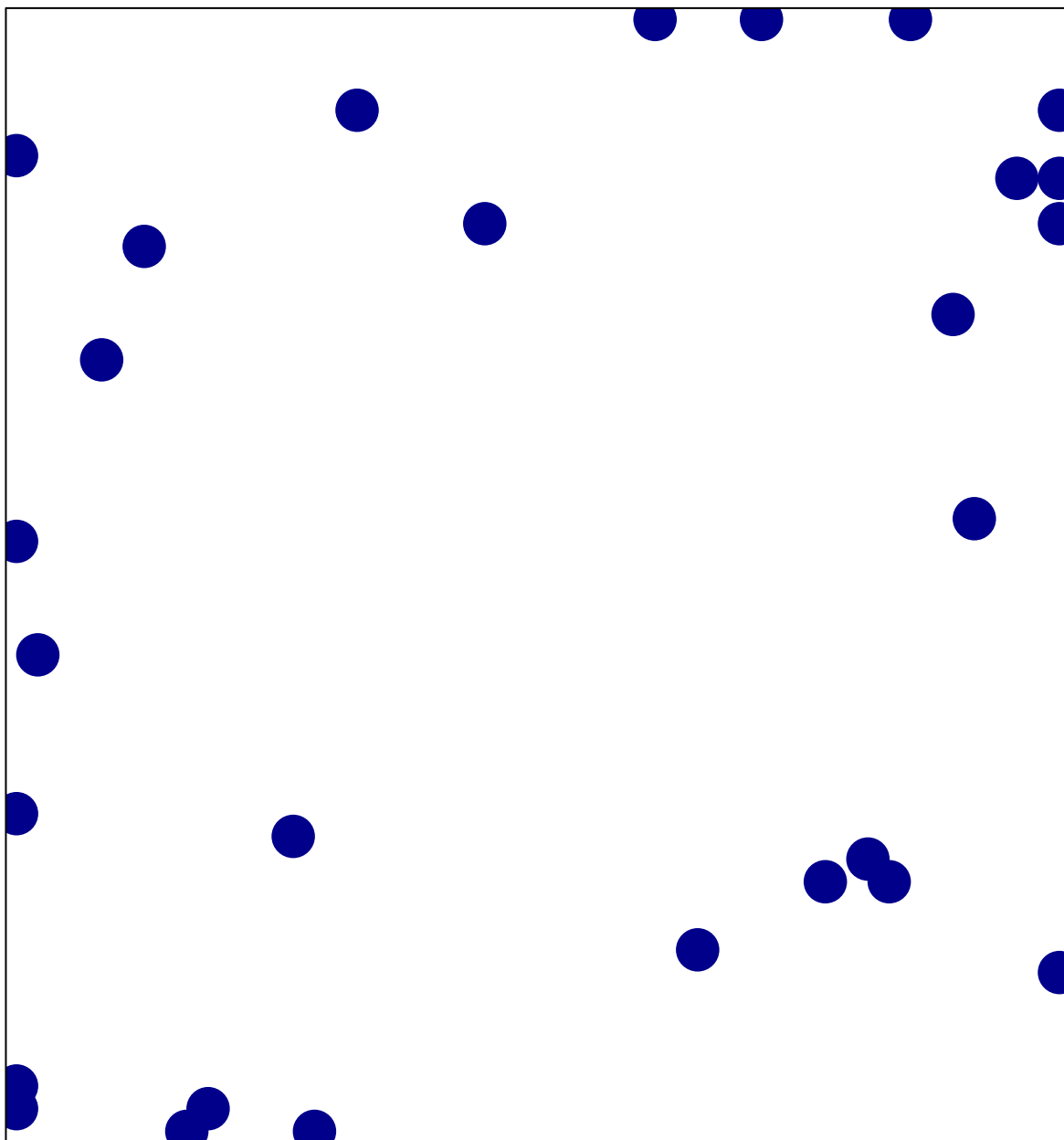
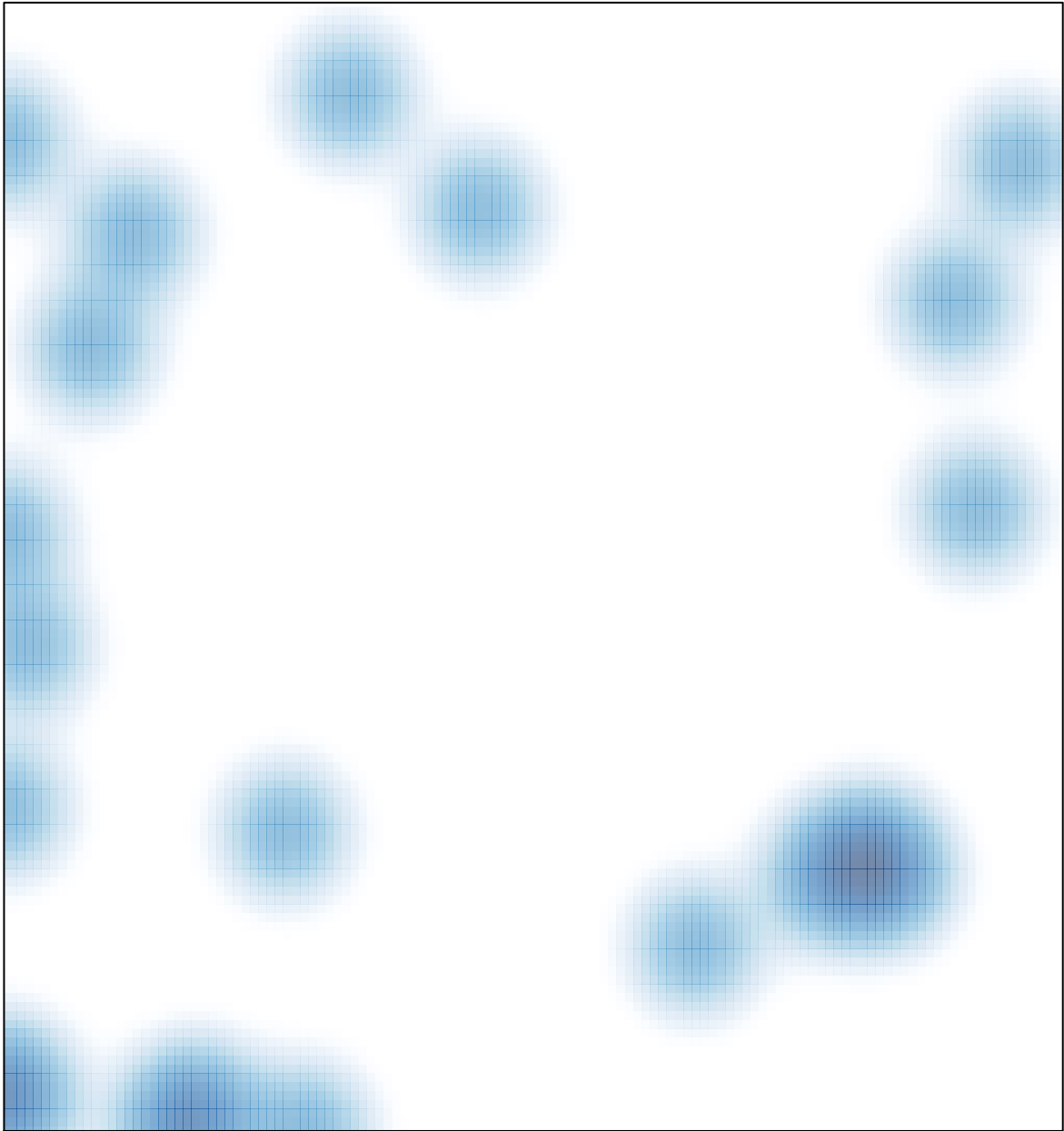


EACTOME_FORMATION_OF_TRANSCRIPTION_COUPLED_NER_TC_NER_REPAIR_COMP



features = 28
chi-square p = 0.82

TION_OF_TRANSCRIPTION_COUPLED_NER_TC_NE



features = 28 , max = 1