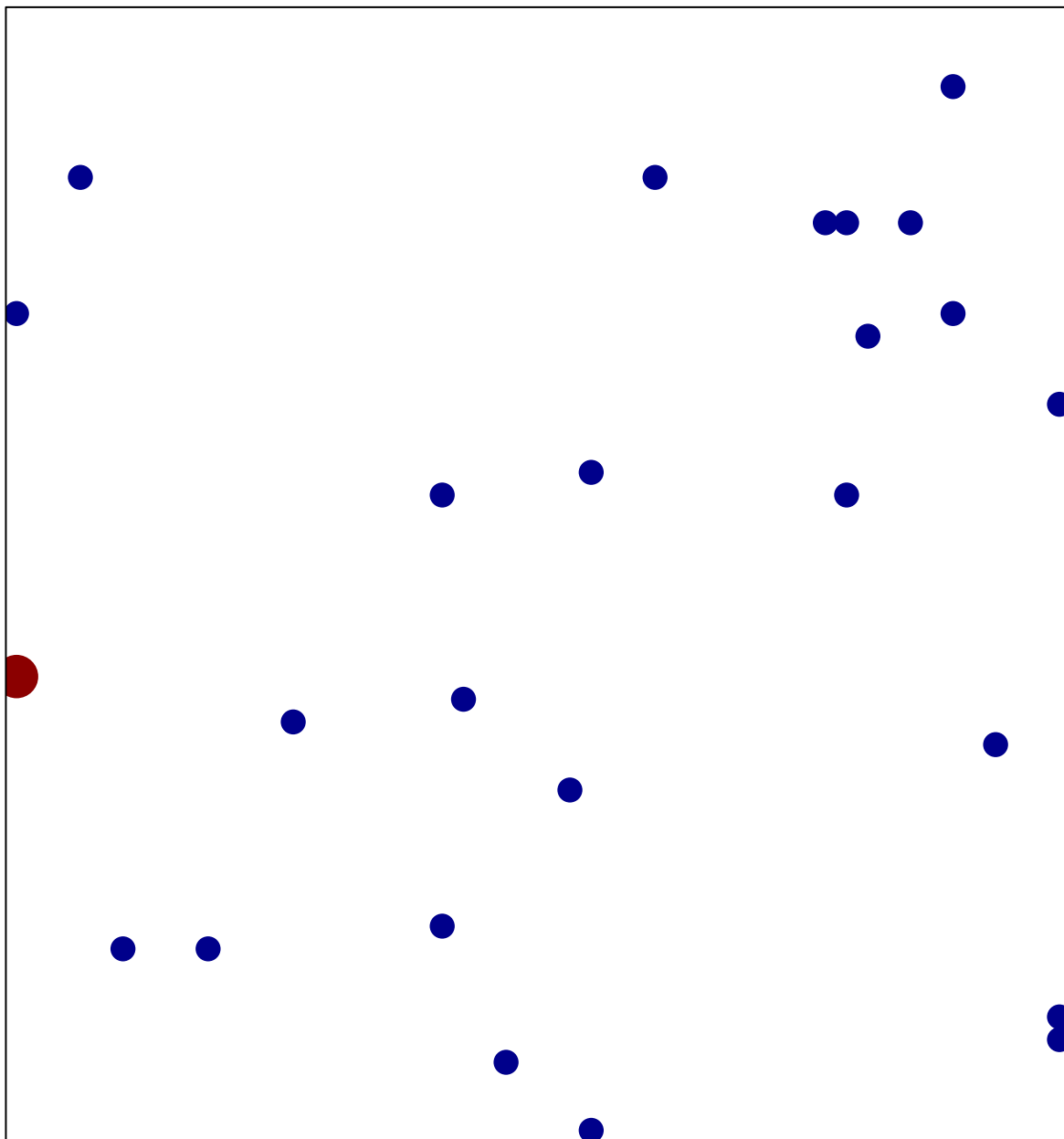
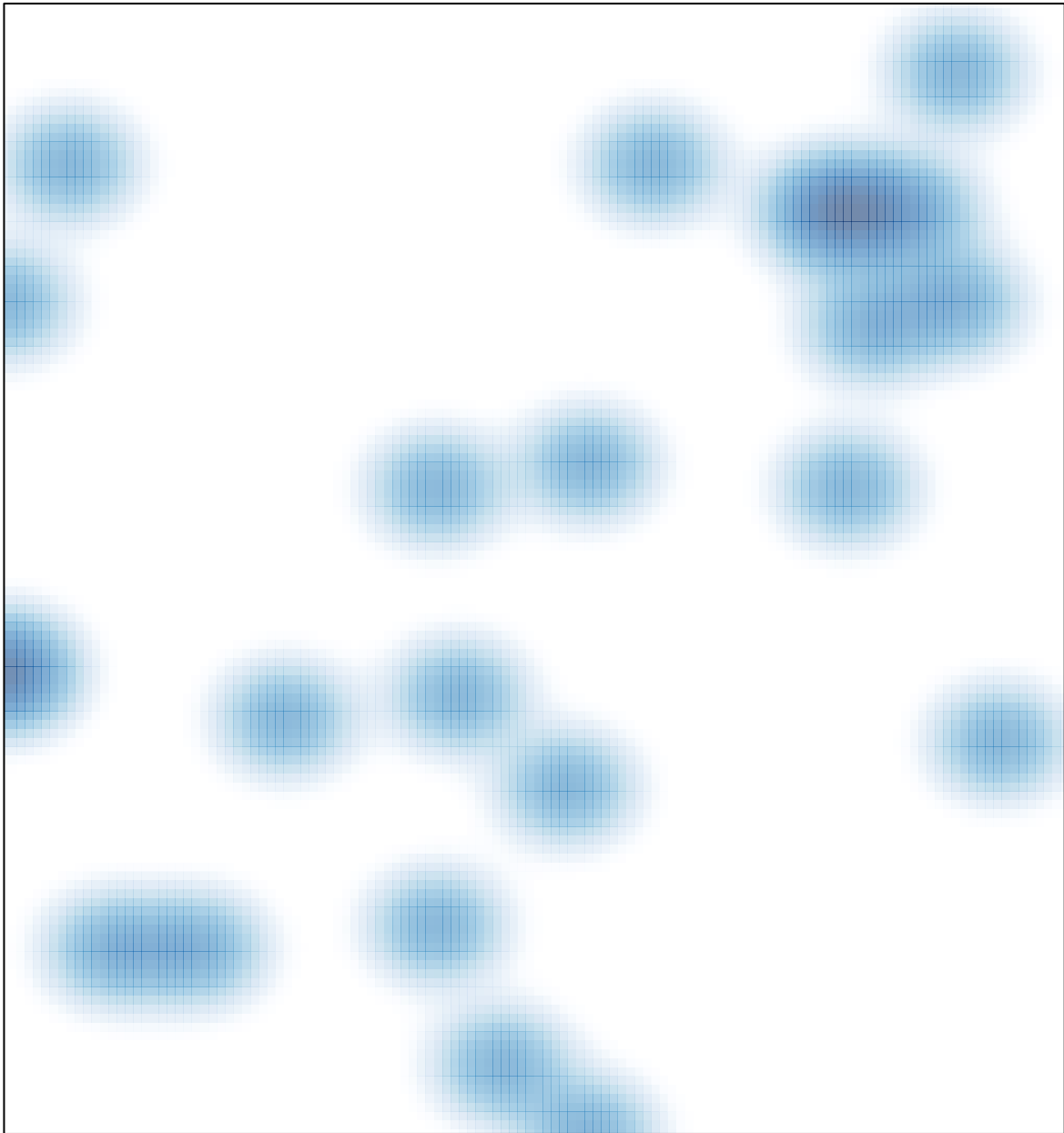


# REACTOME\_NEGATIVE\_REGULATION\_OF\_FGFR\_SIGNALING



# features = 26  
chi-square p = 0.83

# CTOME\_NEGATIVE\_REGULATION\_OF\_FGFR\_SIGNAL



# features = 26 , max = 2