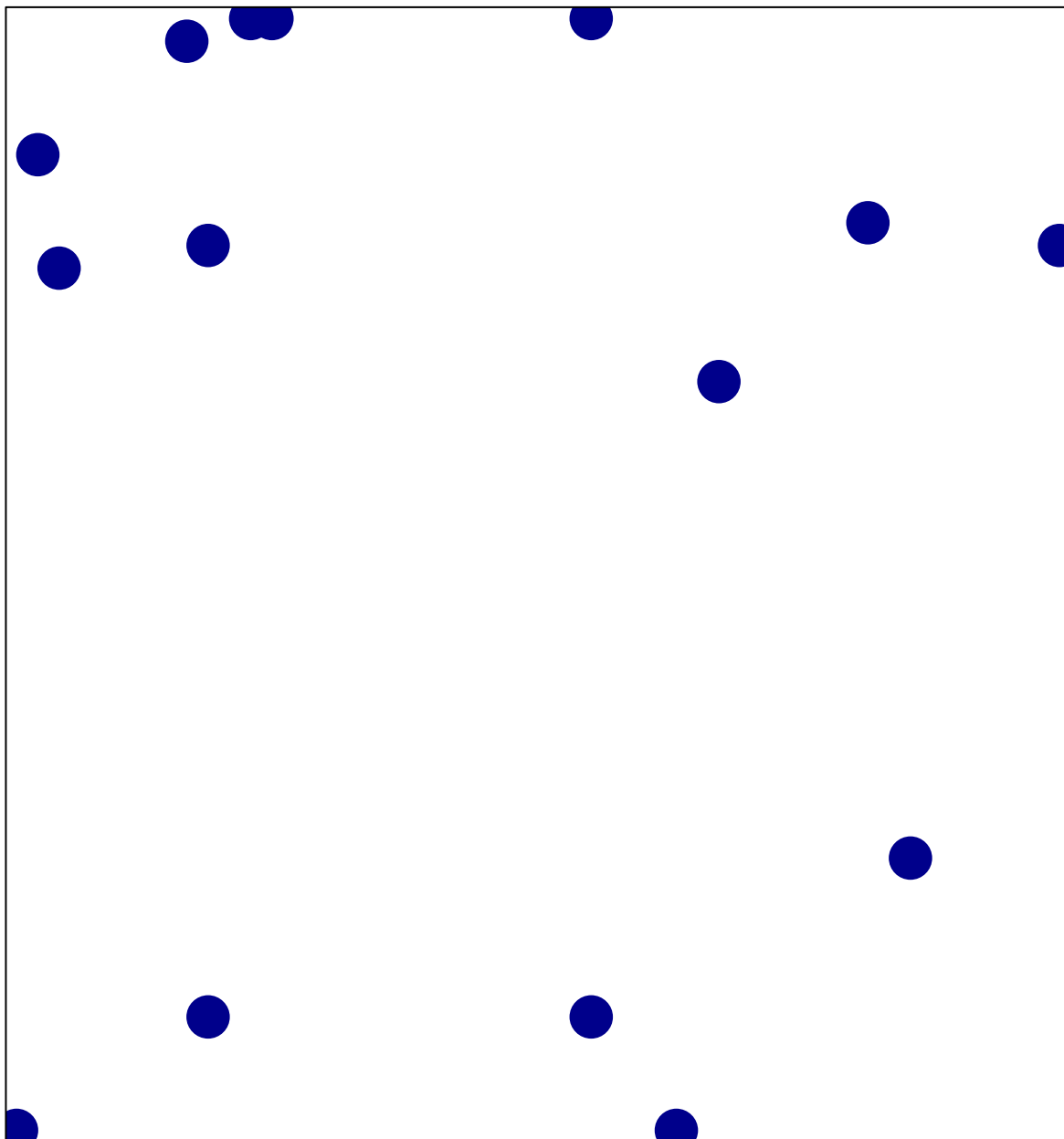
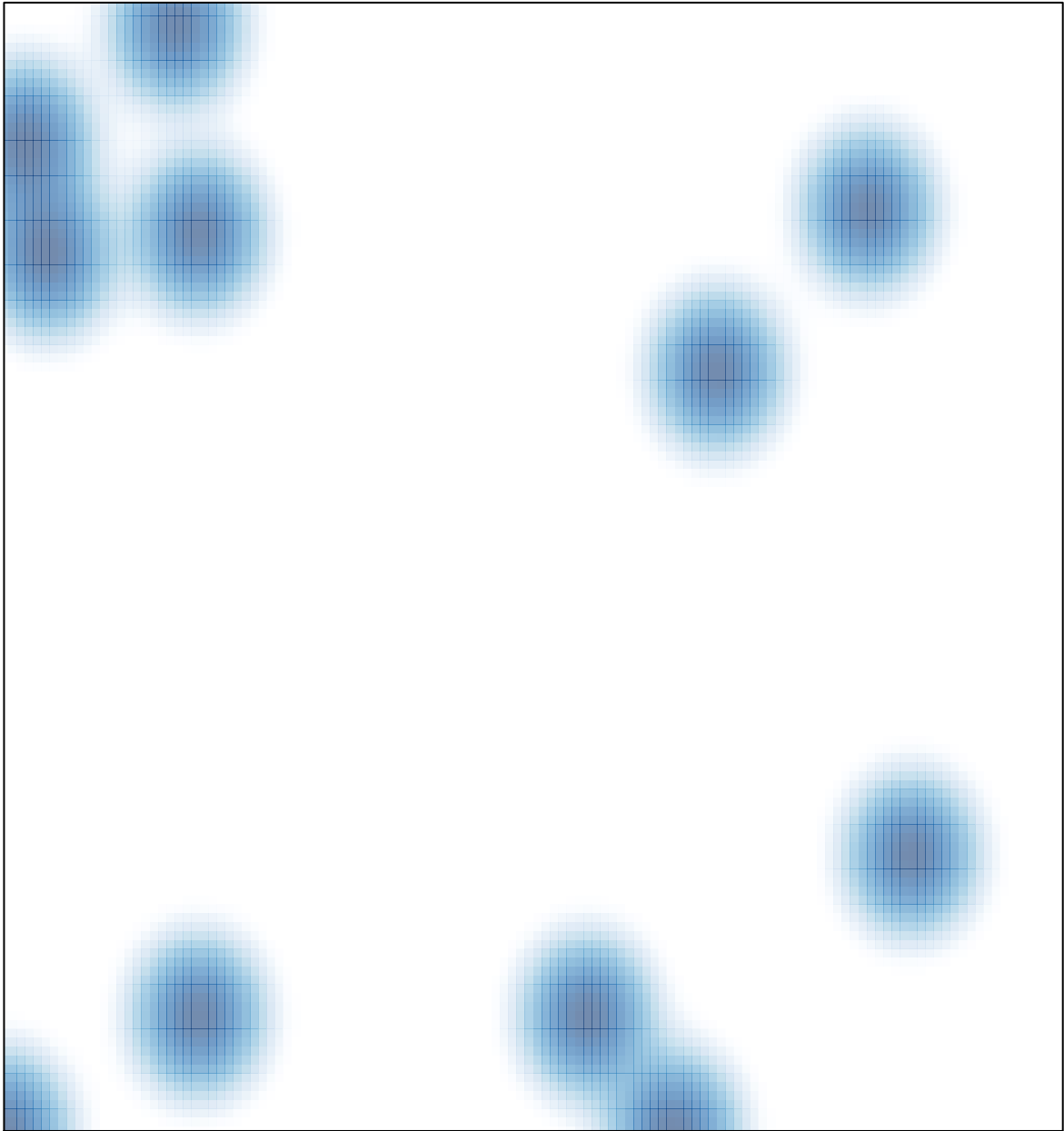


# REACTOME\_GLYCOGEN\_BREAKDOWN\_GLYCOGENOLYSIS



# features = 15  
chi-square p = 0.82

# ACTOME\_GLYCOGEN\_BREAKDOWN\_GLYCOGENOL



# features = 15 , max = 1