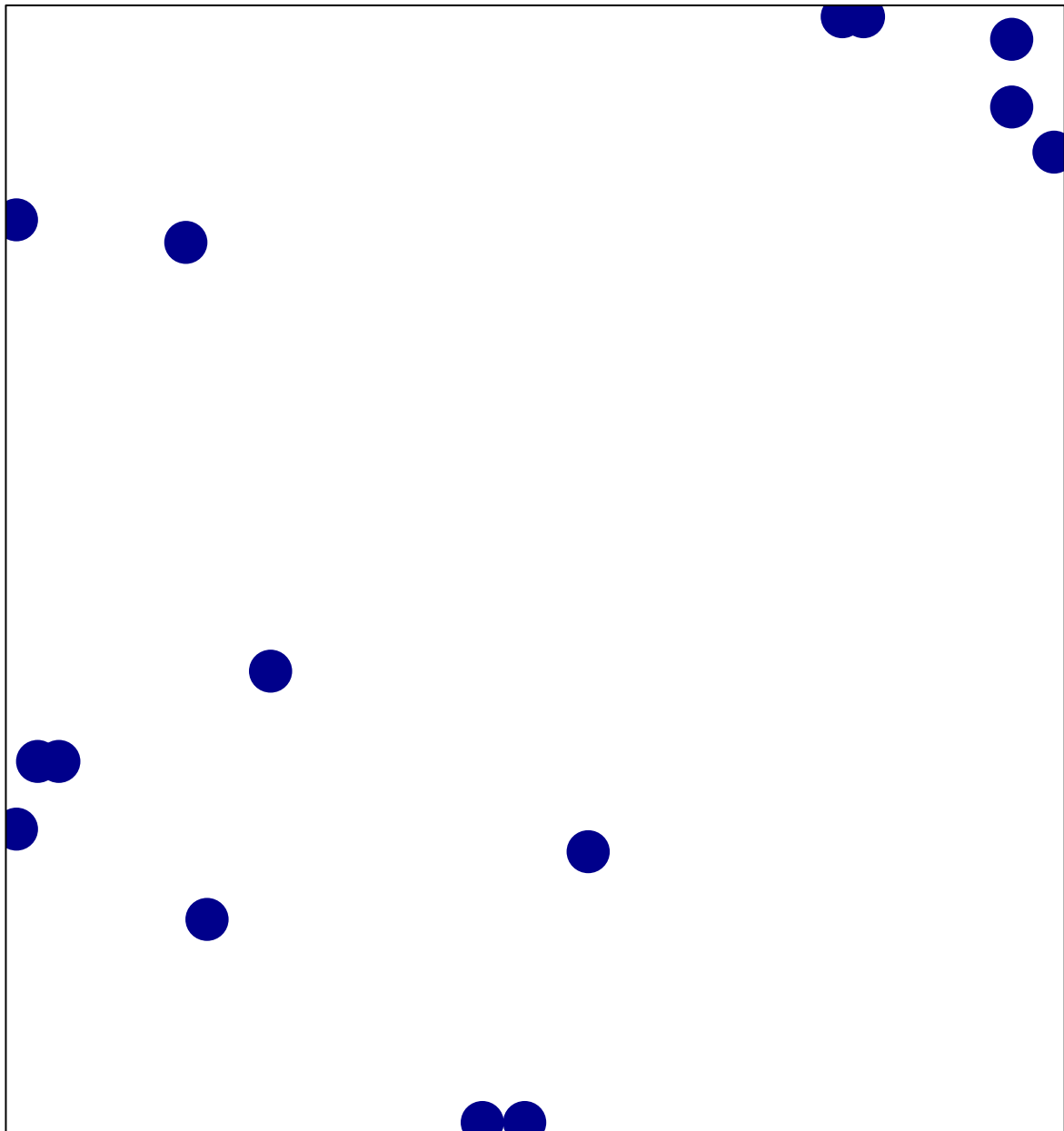


# REACTOME\_GLYCOLYSIS



# features = 15  
chi-square p = 0.84

# REACTOME\_GLUONEOGENESIS



# features = 15 , max = 1