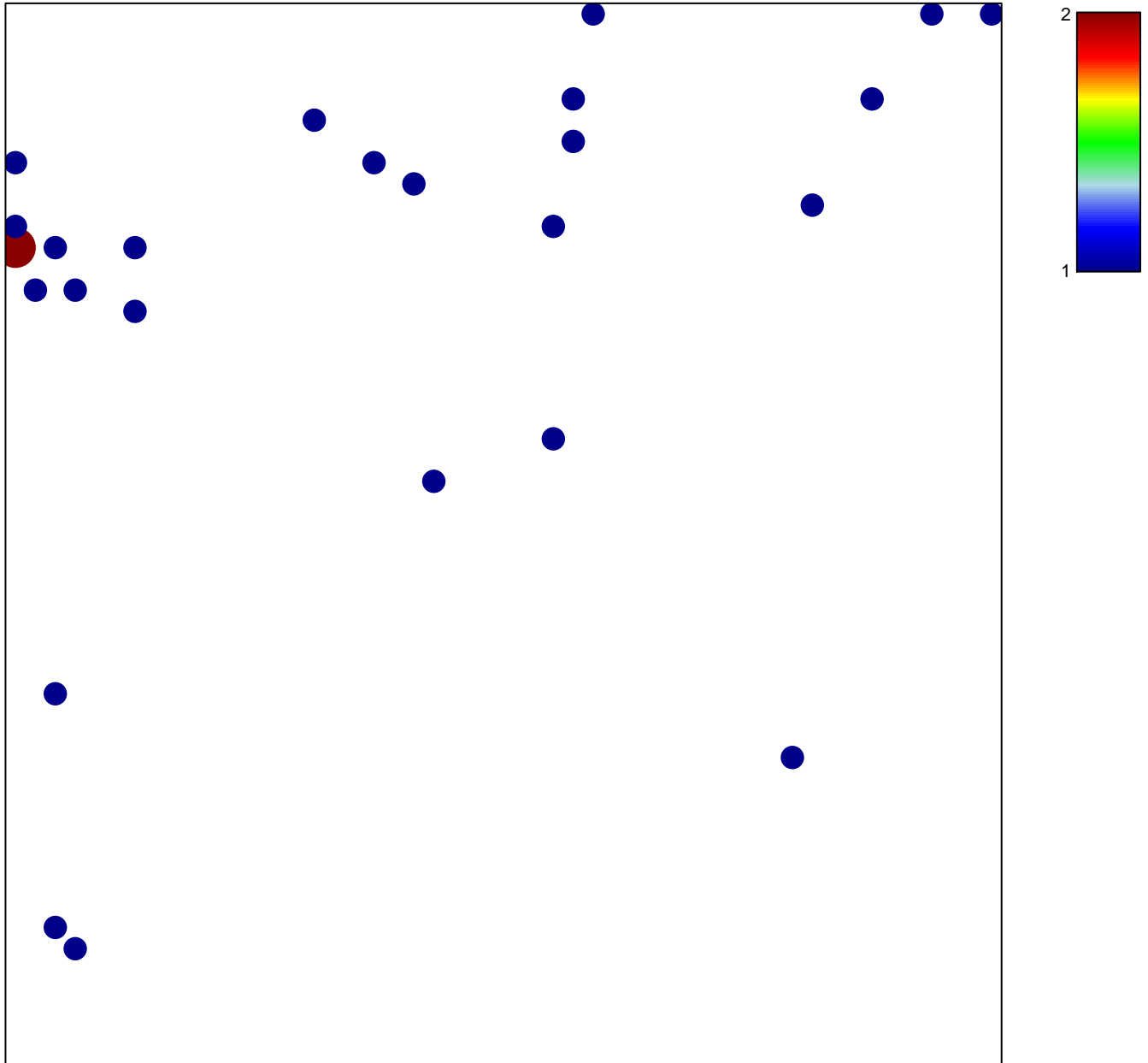


# REACTOME\_GLYCOLYSIS



# features = 26  
chi-square p = 0.81

# REACTOME\_GLYCOGENOLYSIS



# features = 26 , max = 2