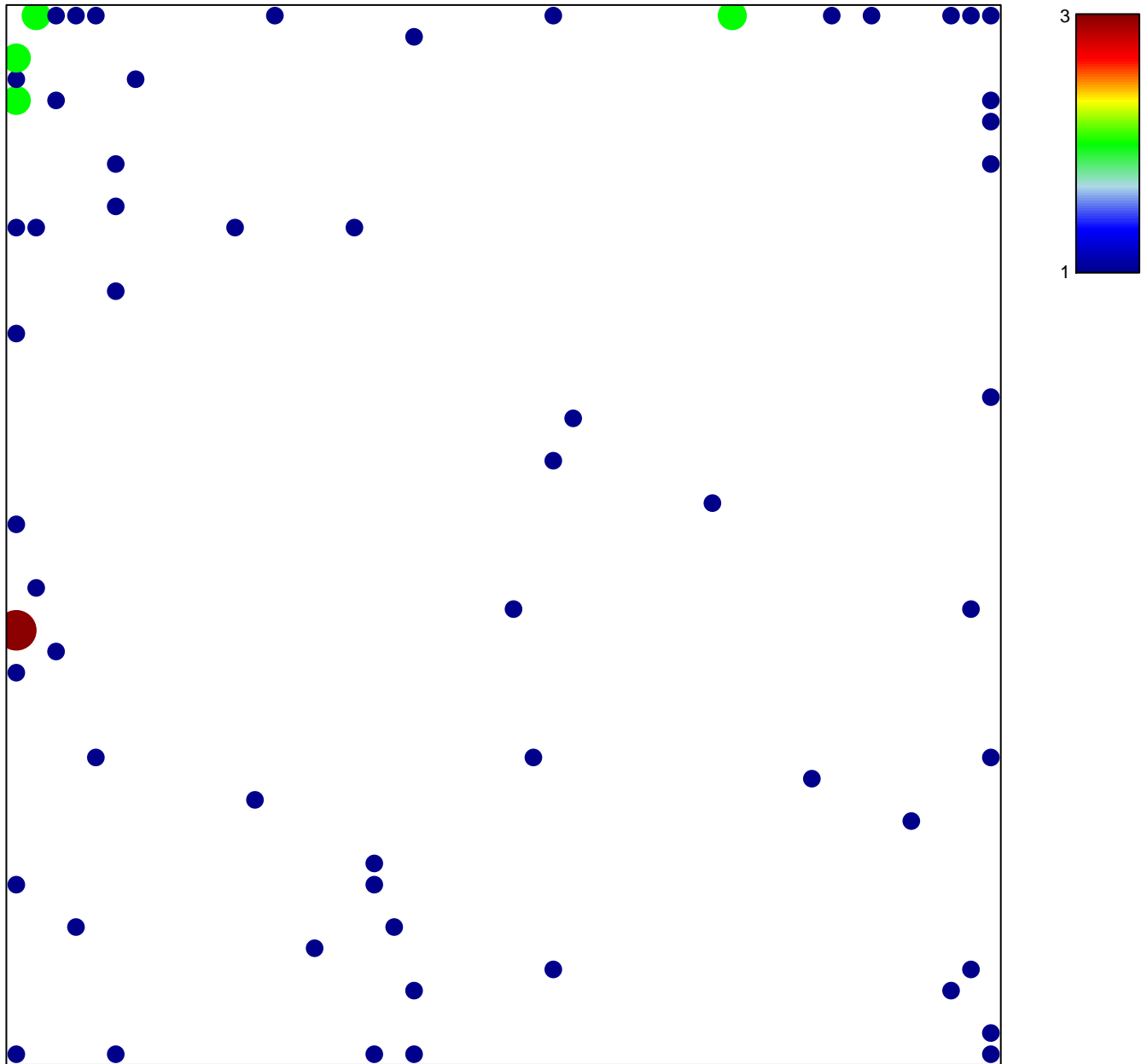


# REACTOME\_METABOLISM\_OF\_NUCLEOTIDES



# features = 68  
chi-square p = 0.7

# REACTOME\_METABOLISM\_OF\_NUCLEOTIDES



# features = 68 , max = 3