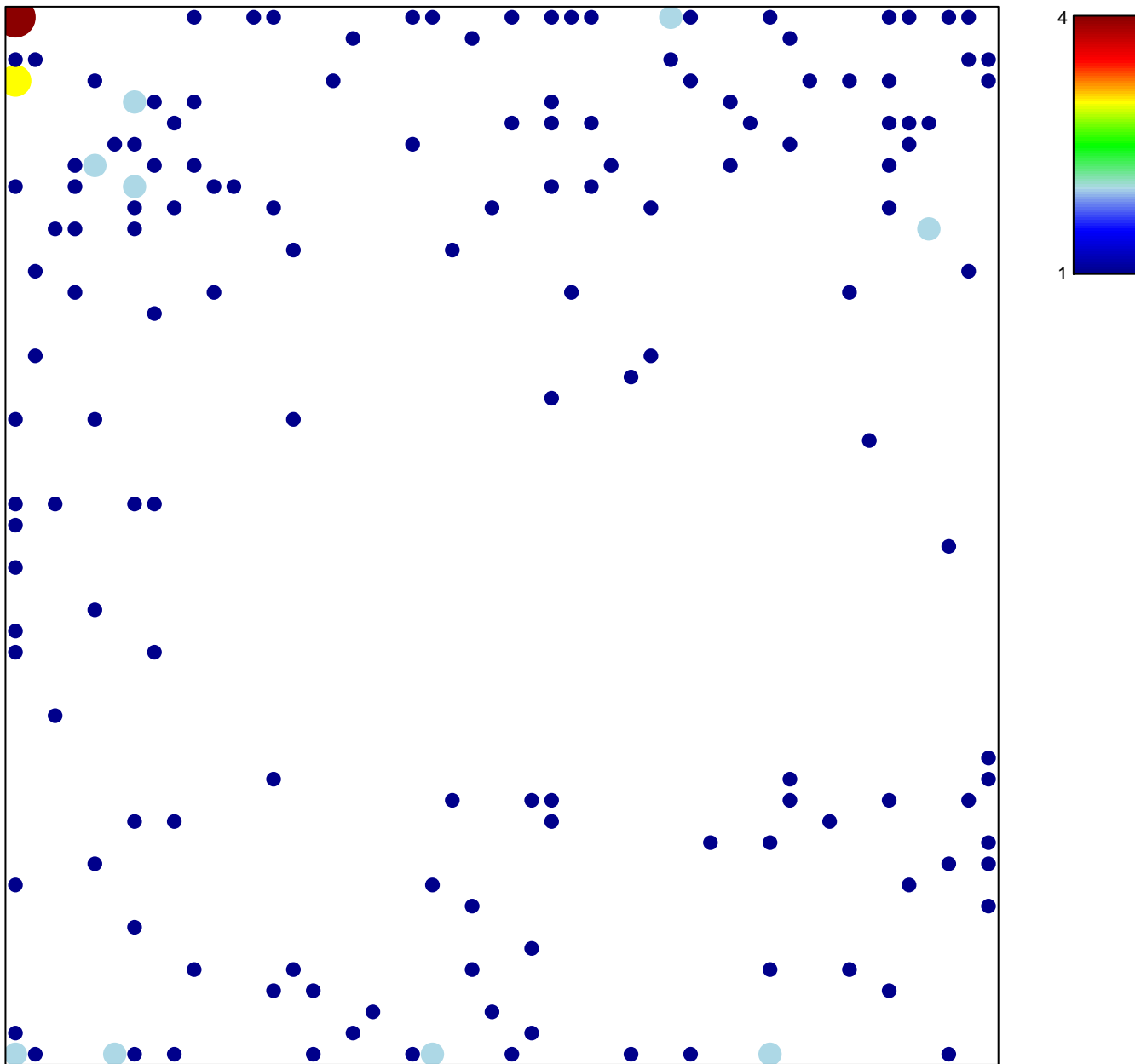
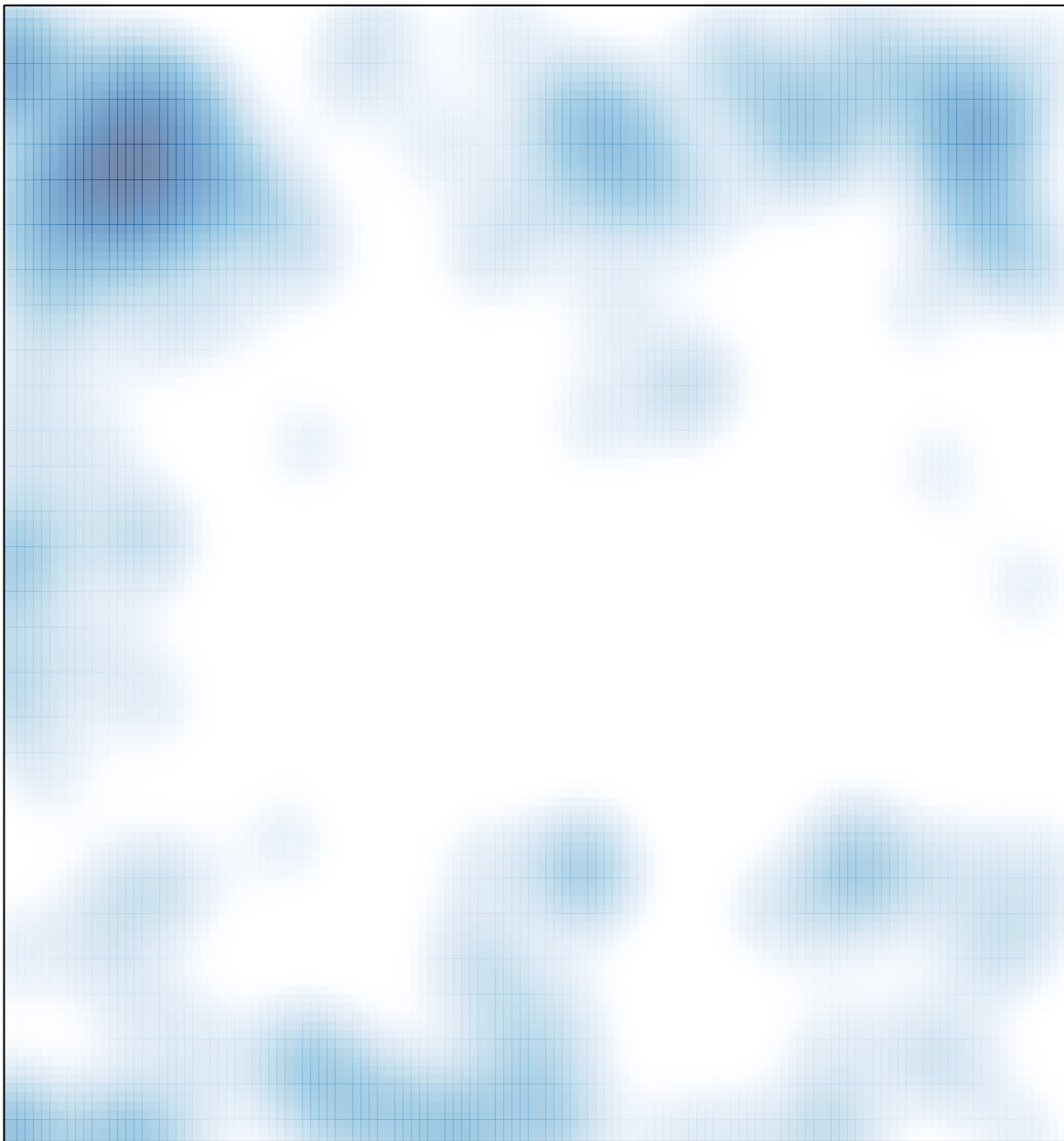


# REACTOME\_METABOLISM\_OF\_RNA



# features = 171  
chi-square p = 0.7

# REACTOME\_METABOLISM\_OF\_RNA



# features = 171 , max = 4