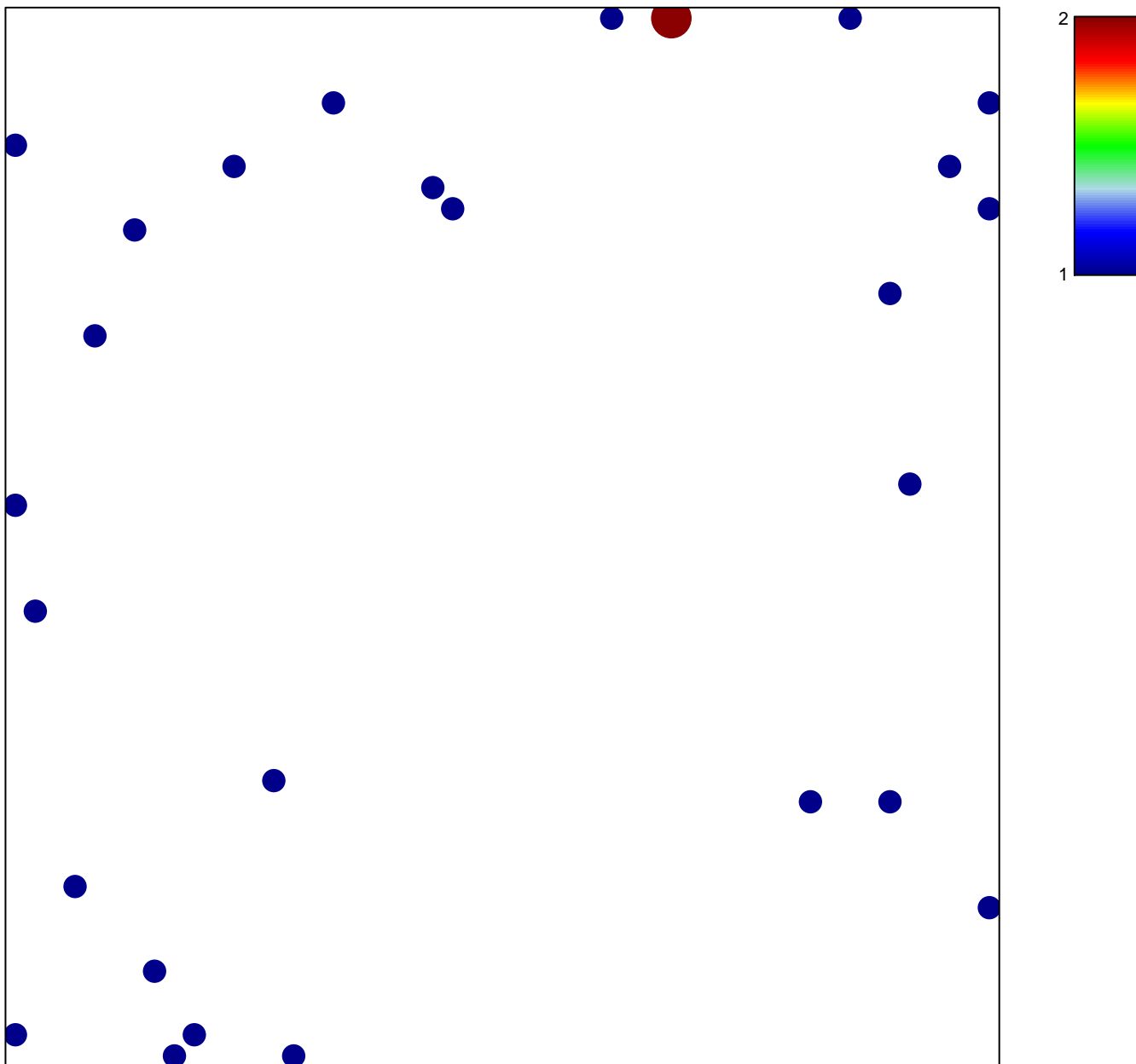
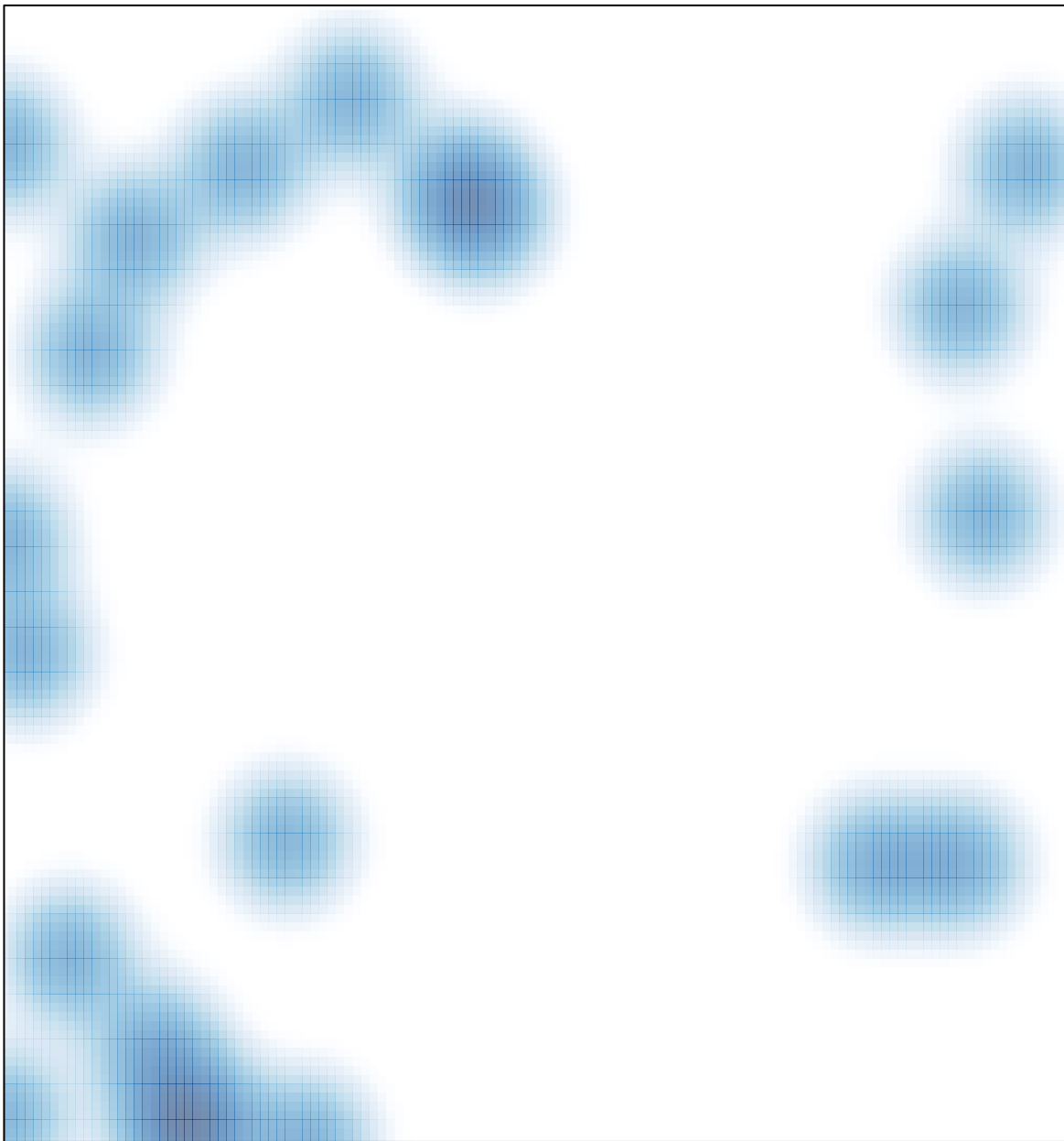


# REACTOME\_MRNA\_CAPPING



# features = 28  
chi-square p = 0.83

# REACTOME\_MRNA\_CAPPING



# features = 28 , max = 2