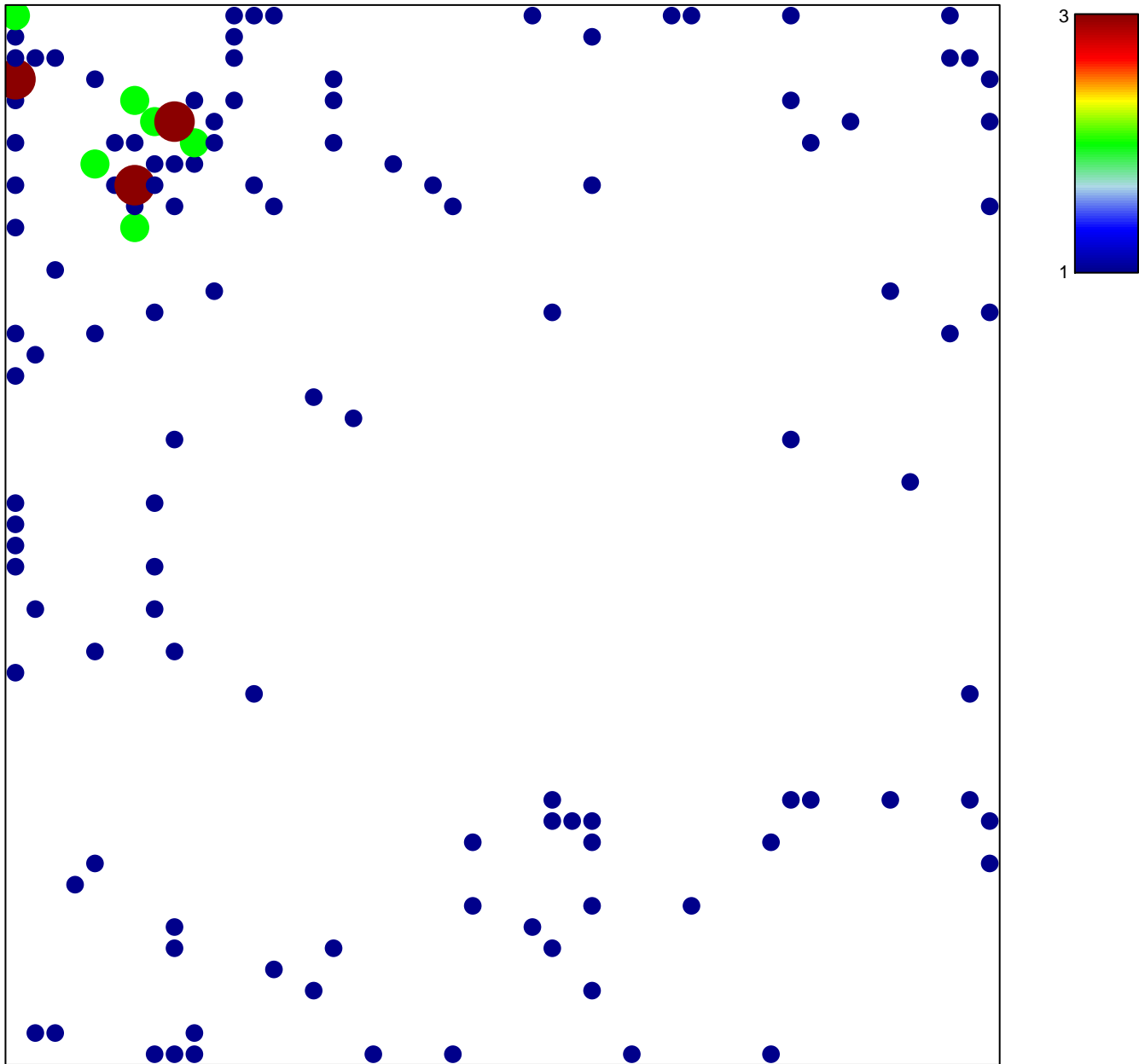
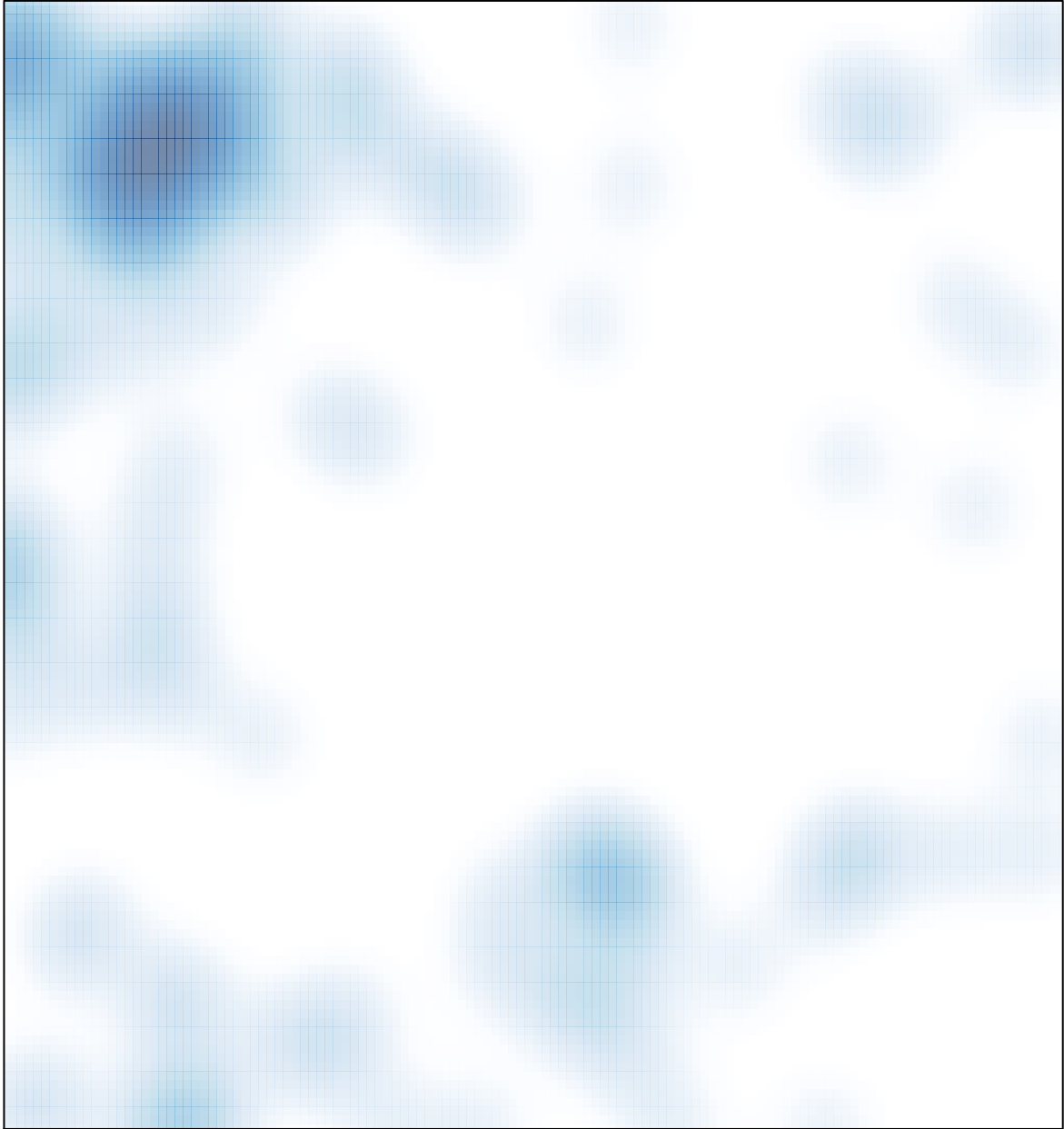


# REACTOME\_PROCESSING\_OF\_CAPPED\_INTRON\_CONTAINING\_PRE\_MRNA



# features = 136  
chi-square p = 0.72

# \_PROCESSING\_OF\_CAPPED\_INTRON\_CONTAINING



# features = 136 , max = 3