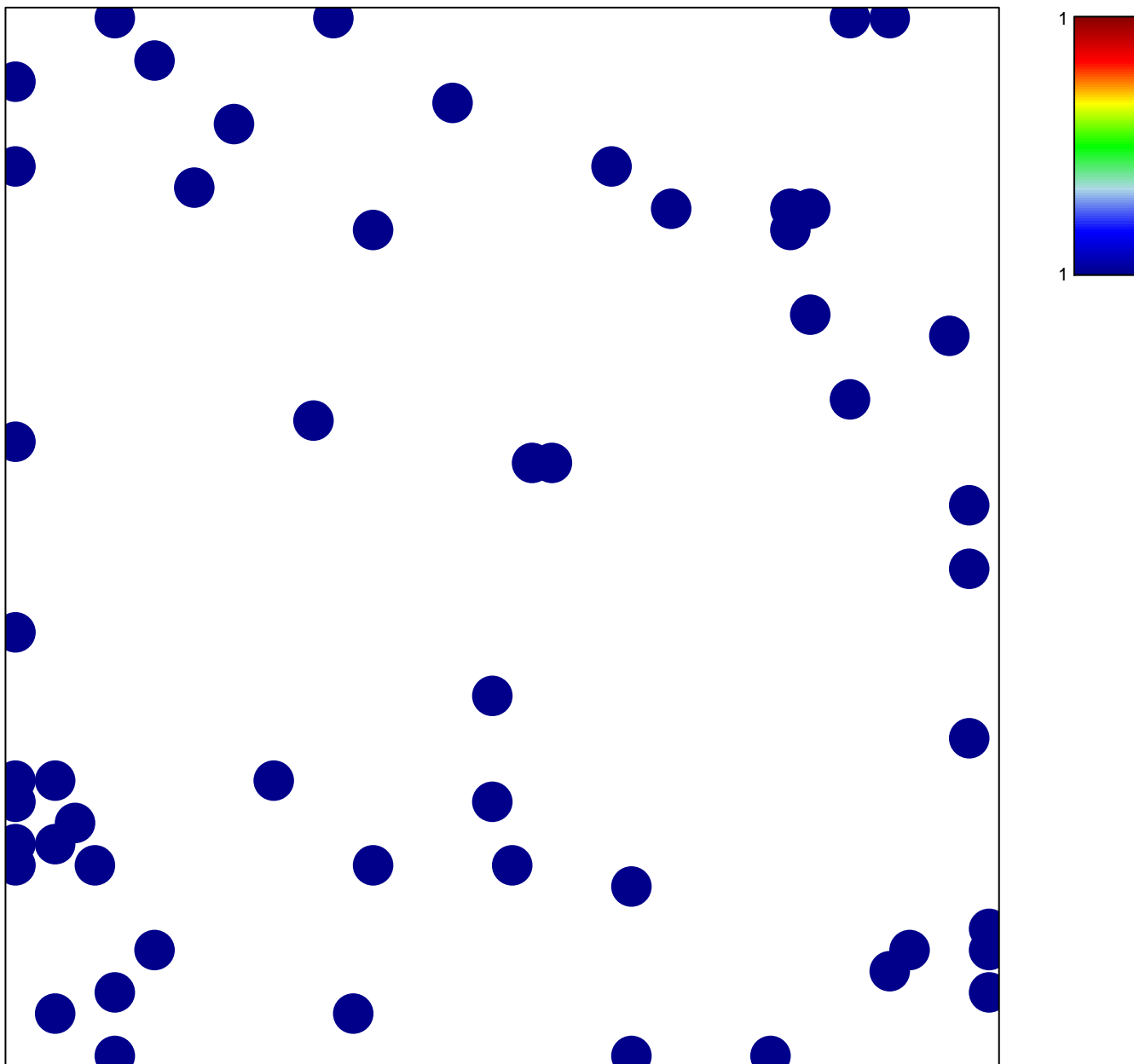
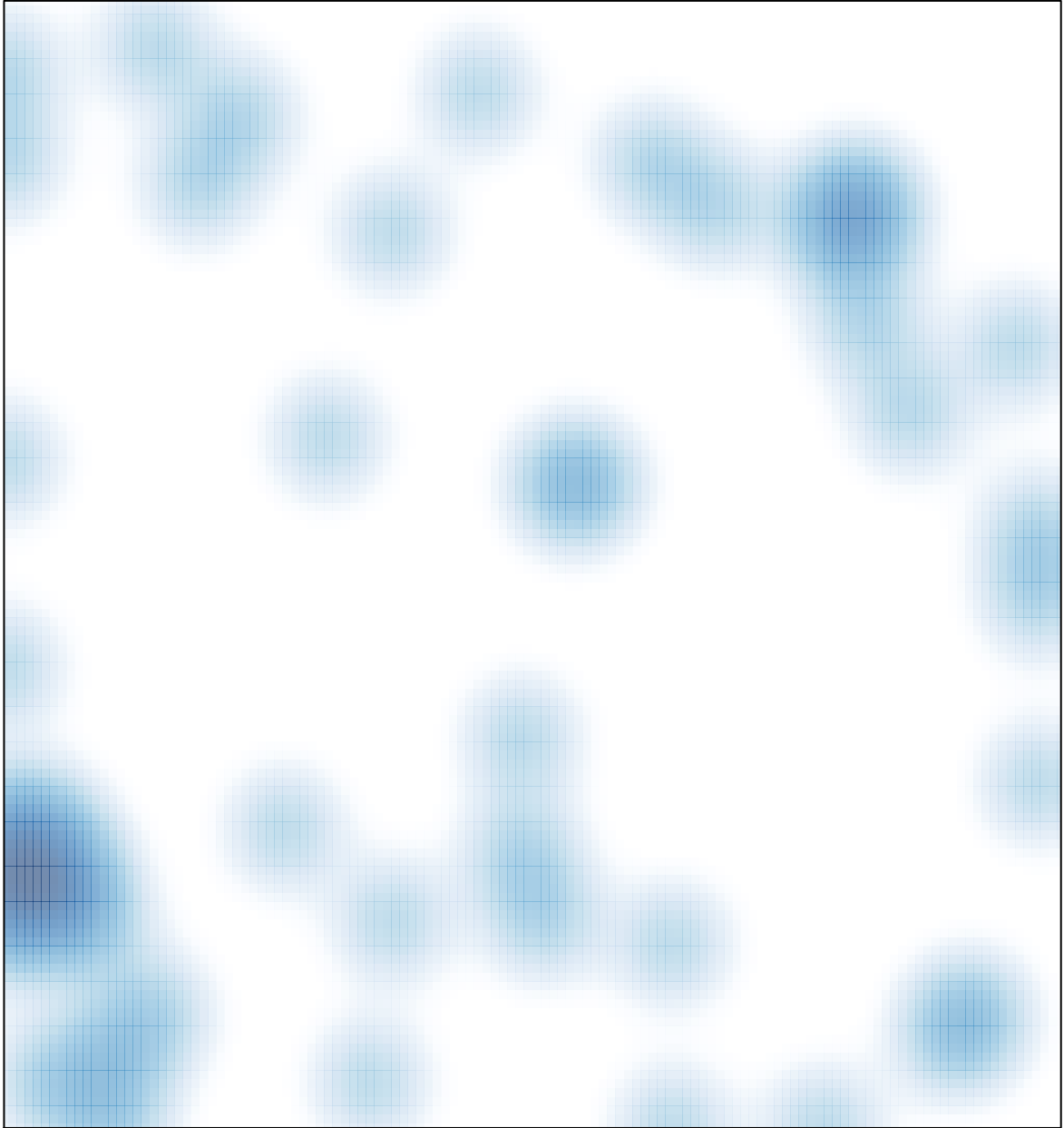


# REACTOME\_CELL\_DEATH\_SIGNALLING\_VIA\_NRAGE\_NRIF\_AND\_NAID



# features = 53  
chi-square p = 0.81

# ME\_CELL\_DEATH\_SIGNALLING\_VIA\_NRAGE\_NRIF\_A



# features = 53 , max = 1