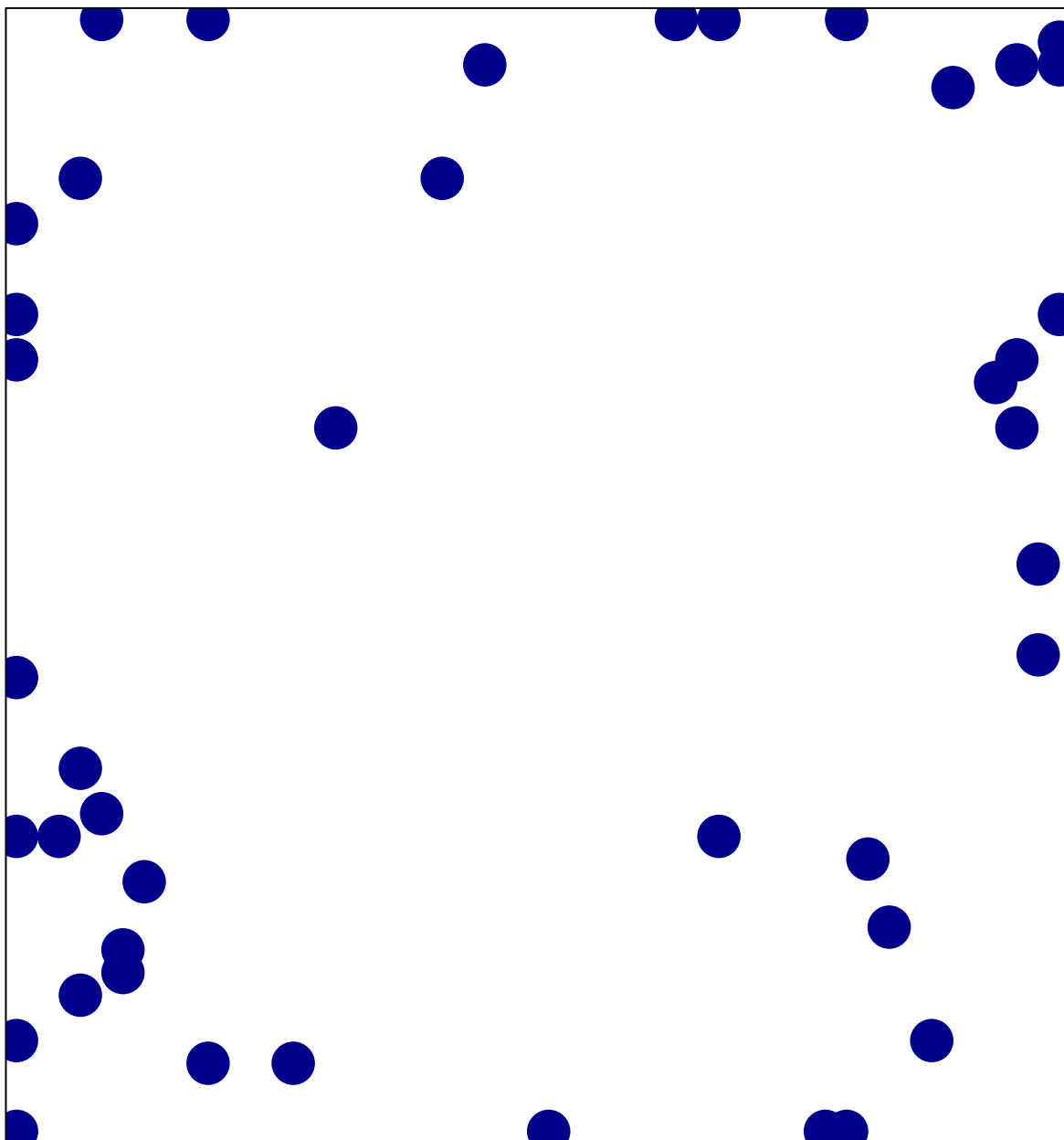
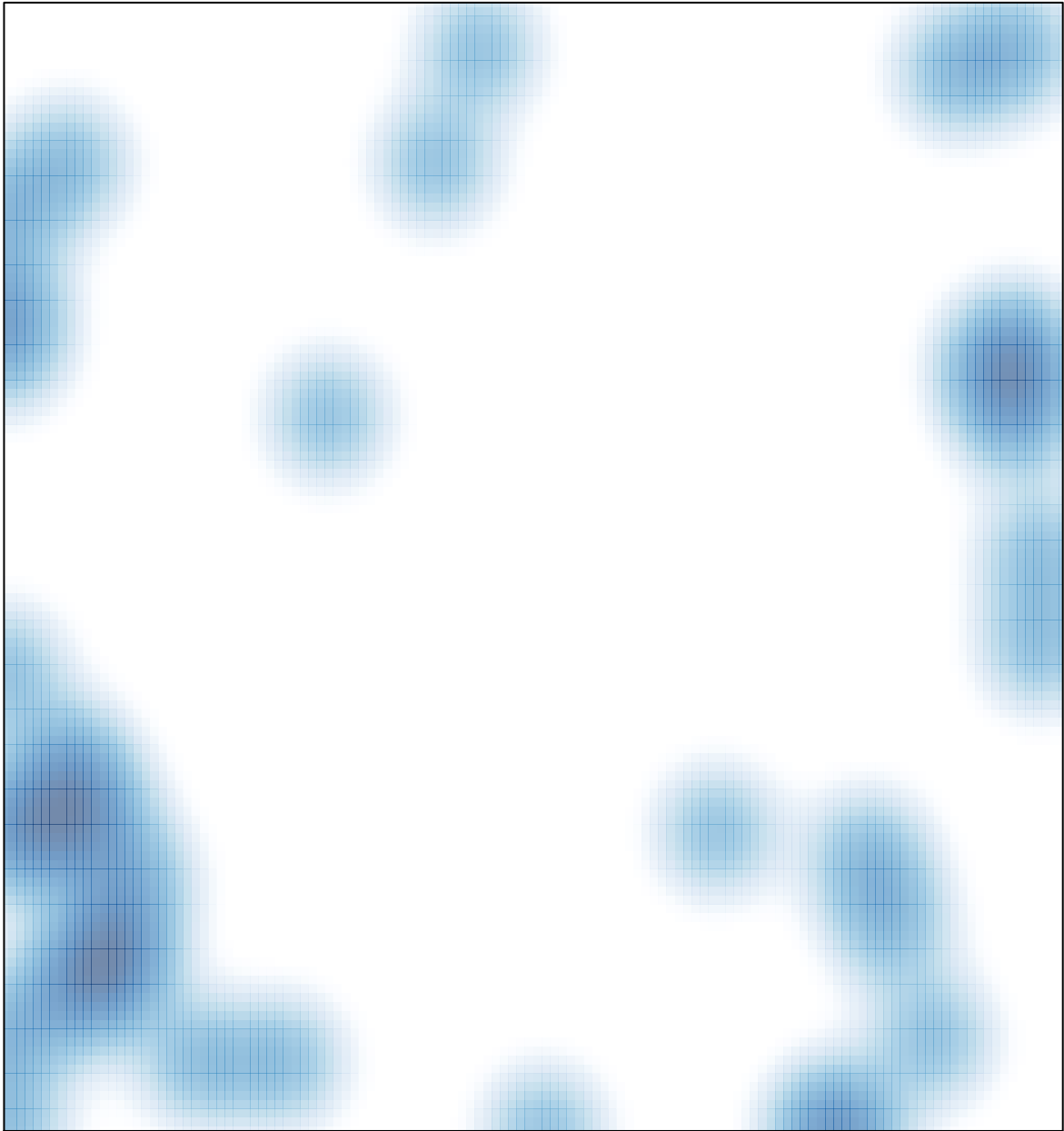


# REACTOME\_MAP\_KINASE\_ACTIVATION\_IN\_TLR\_CASCADE



# features = 42  
chi-square p = 0.8

# ACTOME\_MAP\_KINASE\_ACTIVATION\_IN\_TLR\_CASE



# features = 42 , max = 1