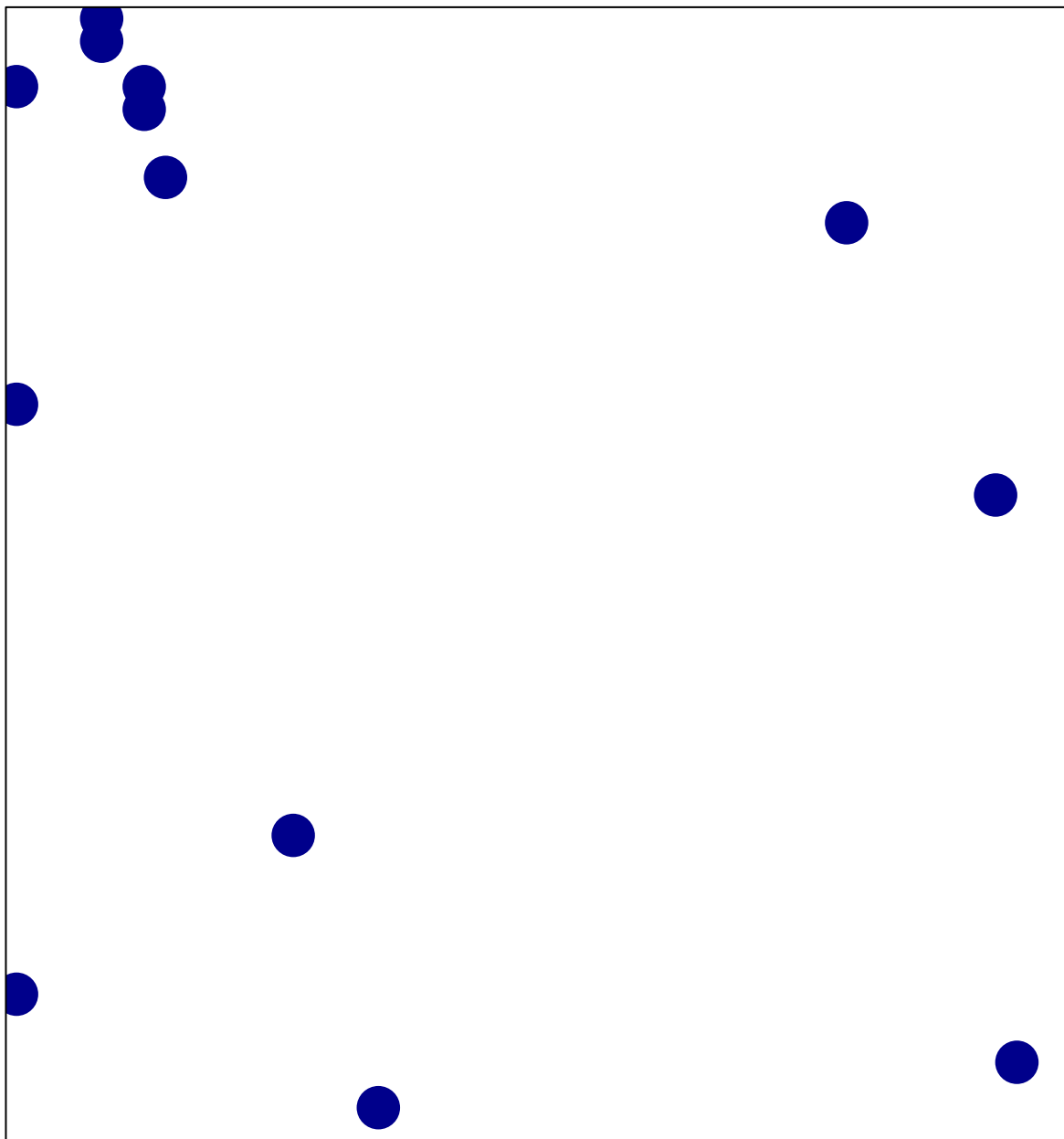
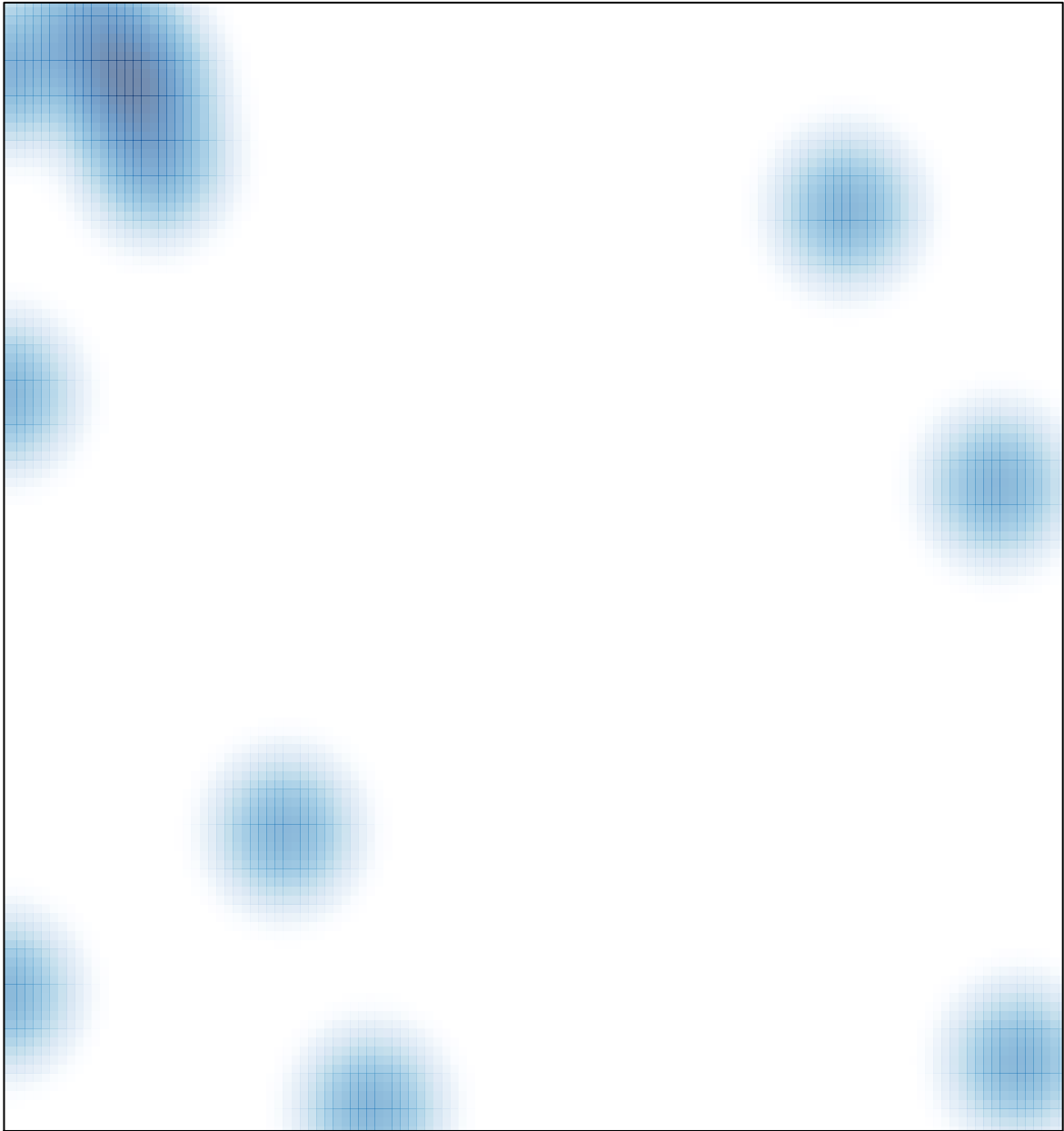


# REACTOME\_APOPTOSIS\_INDUCED\_DNA\_FRAGMENTATION



# features = 13  
chi-square p = 0.83

# ACTOME\_APOPTOSIS\_INDUCED\_DNA\_FRAGMENTA



# features = 13 , max = 1