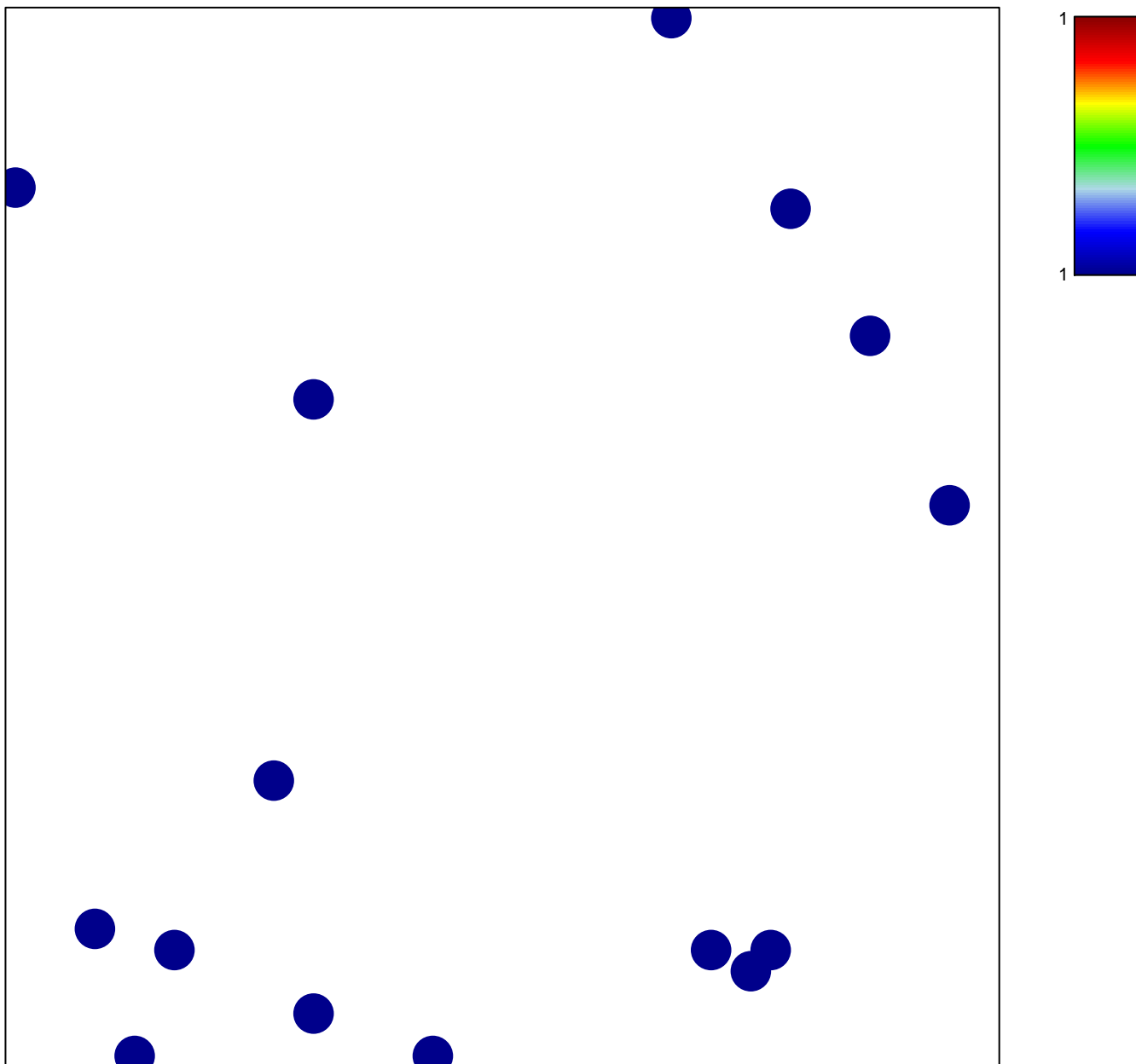
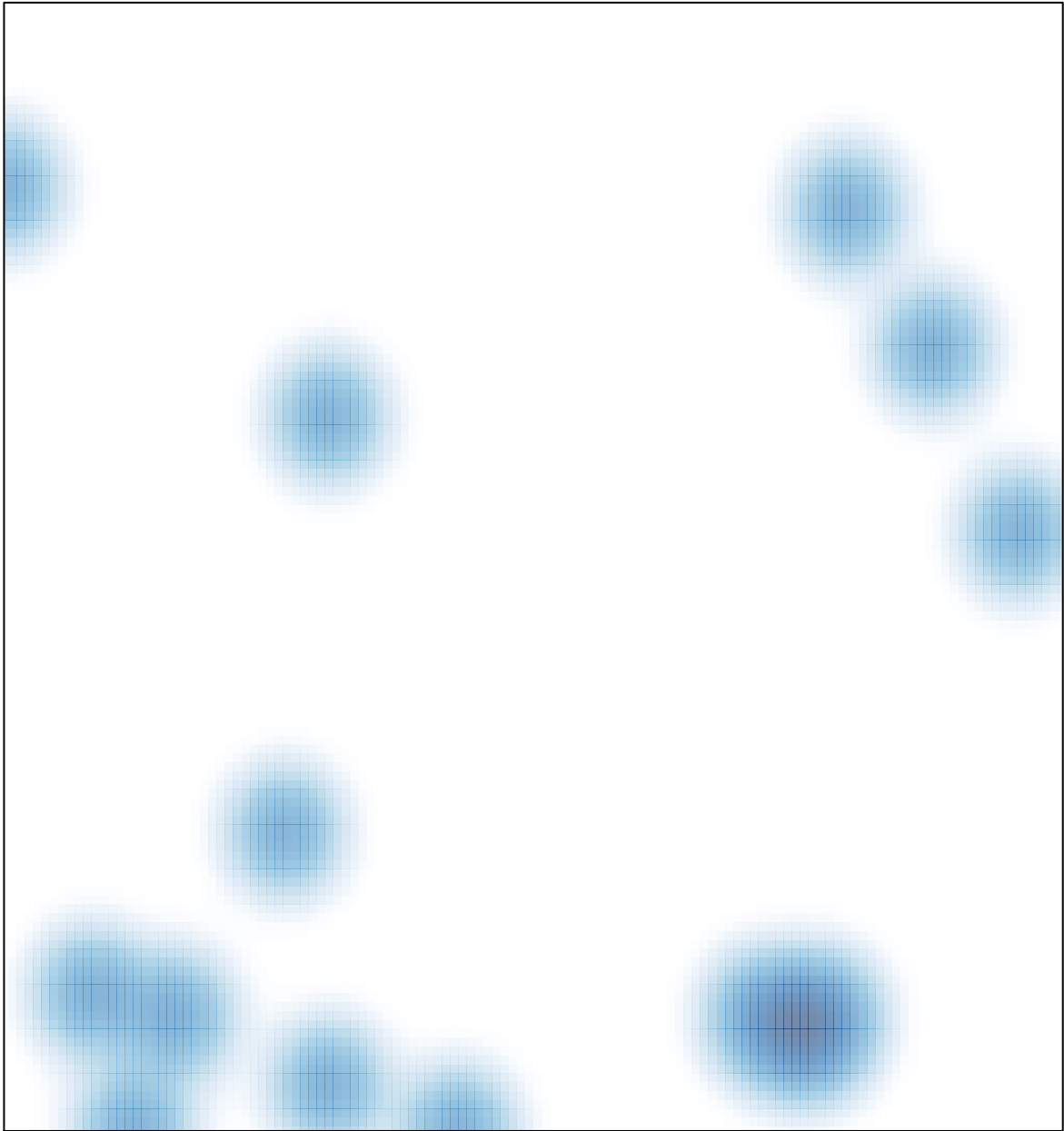


# REACTOME\_PROTEOLYTIC\_CLEAVAGE\_OF\_SNARE\_COMPLEX\_PROTEINS



# features = 15  
chi-square p = 0.84

# E\_PROTEOLYTIC\_CLEAVAGE\_OF\_SNARE\_COMPLEX



# features = 15 , max = 1