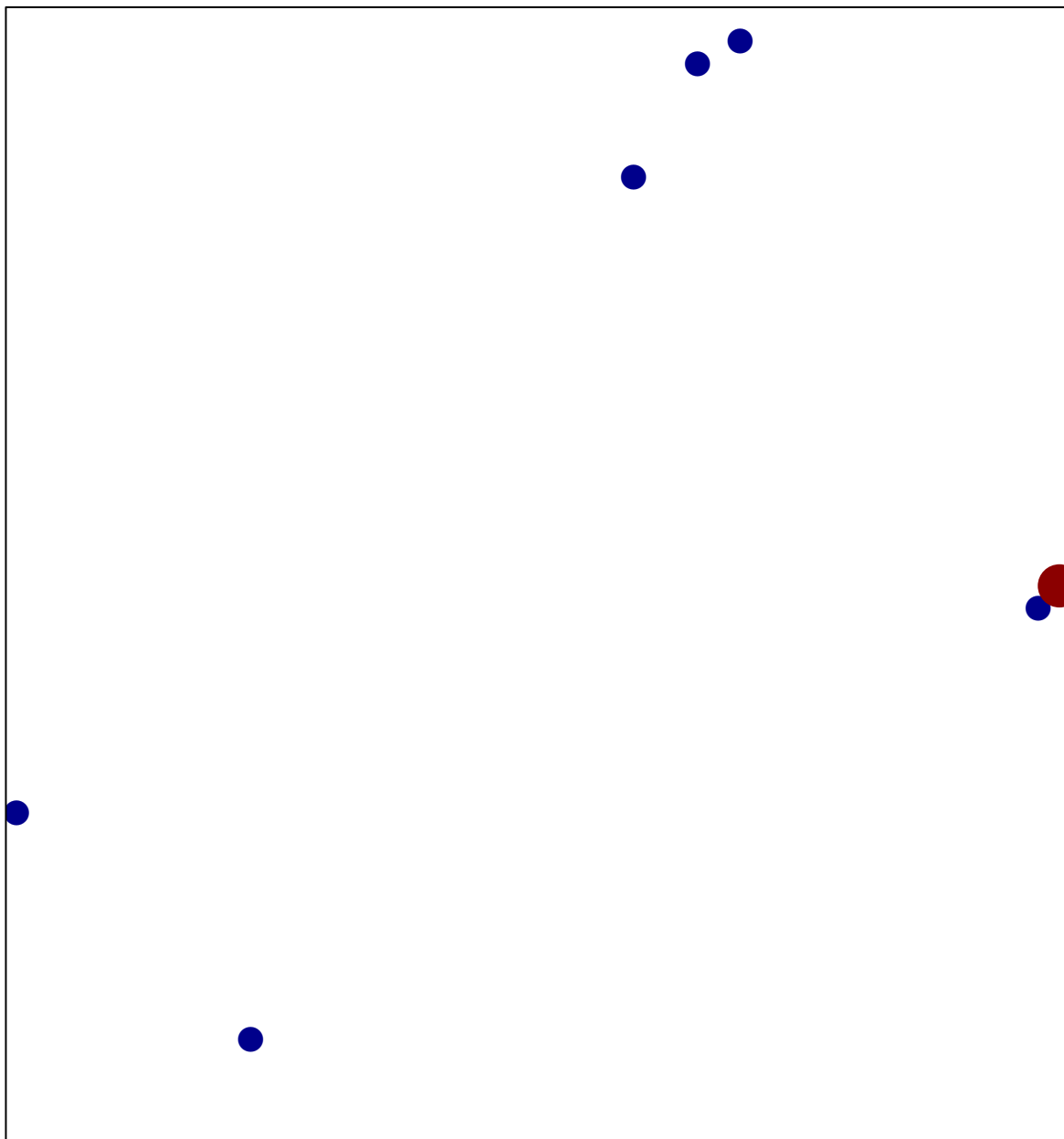
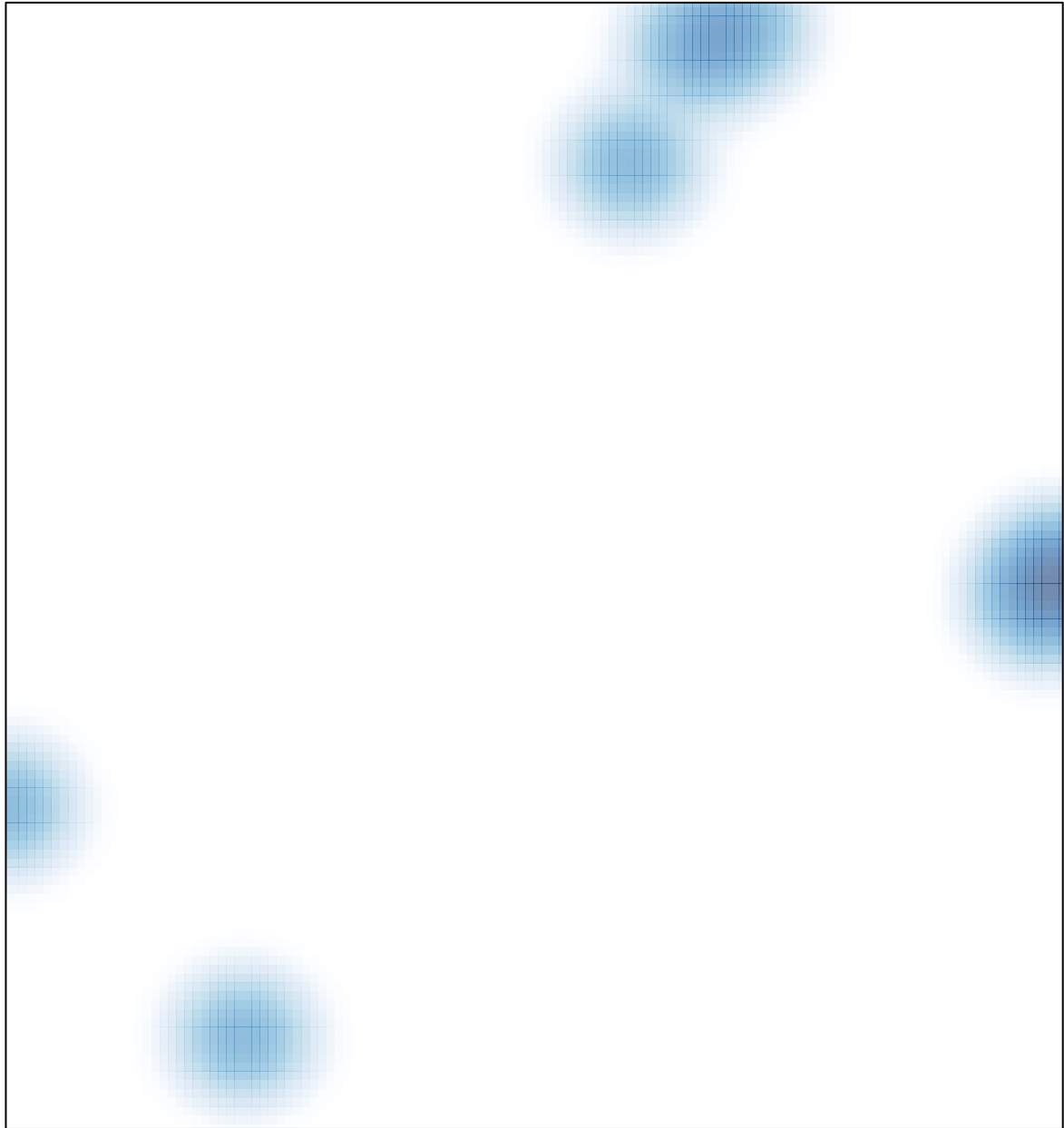


# REACTOME\_METABOLISM\_OF\_PROTEINS



# features = 8  
chi-square p = 0.83

# REACTOME\_METABOLISM\_OF\_PROTEINS



# features = 8 , max = 2