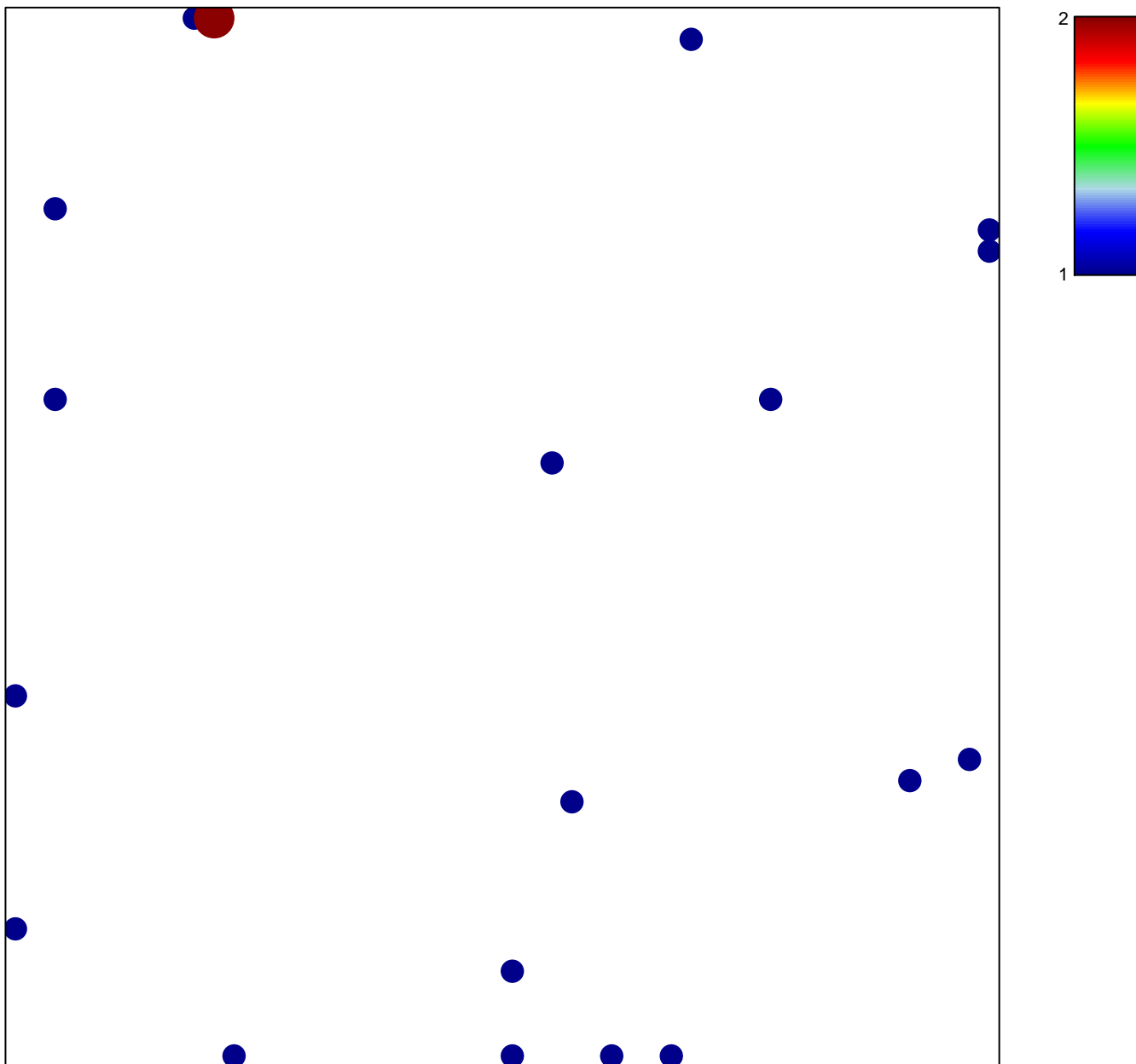
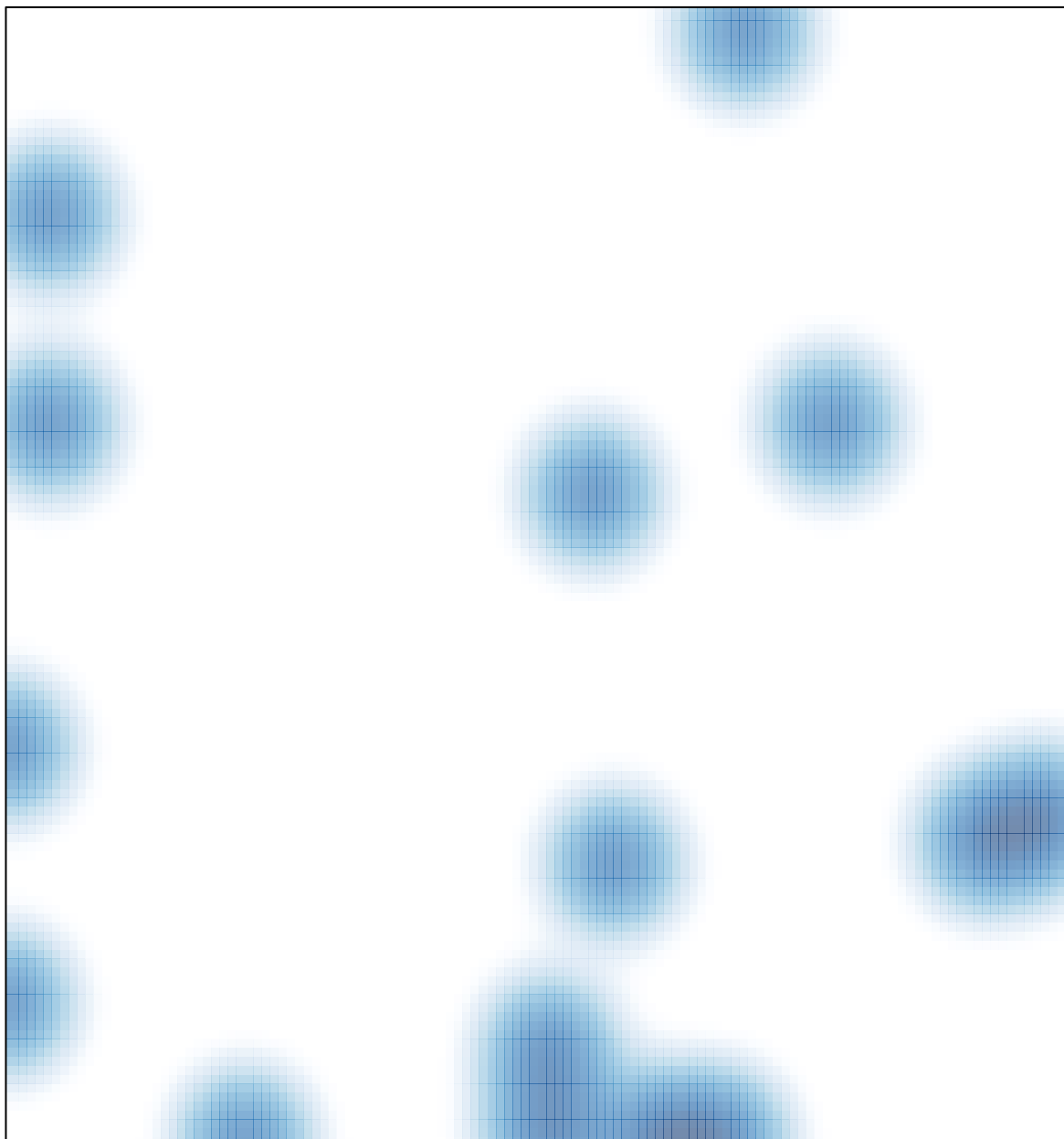


# REACTOME\_PEROXISOMAL\_LIPID\_METABOLISM



# features = 20  
chi-square p = 0.82

# REACTOME\_PEROXISOMAL\_LIPID\_METABOLISM



# features = 20 , max = 2