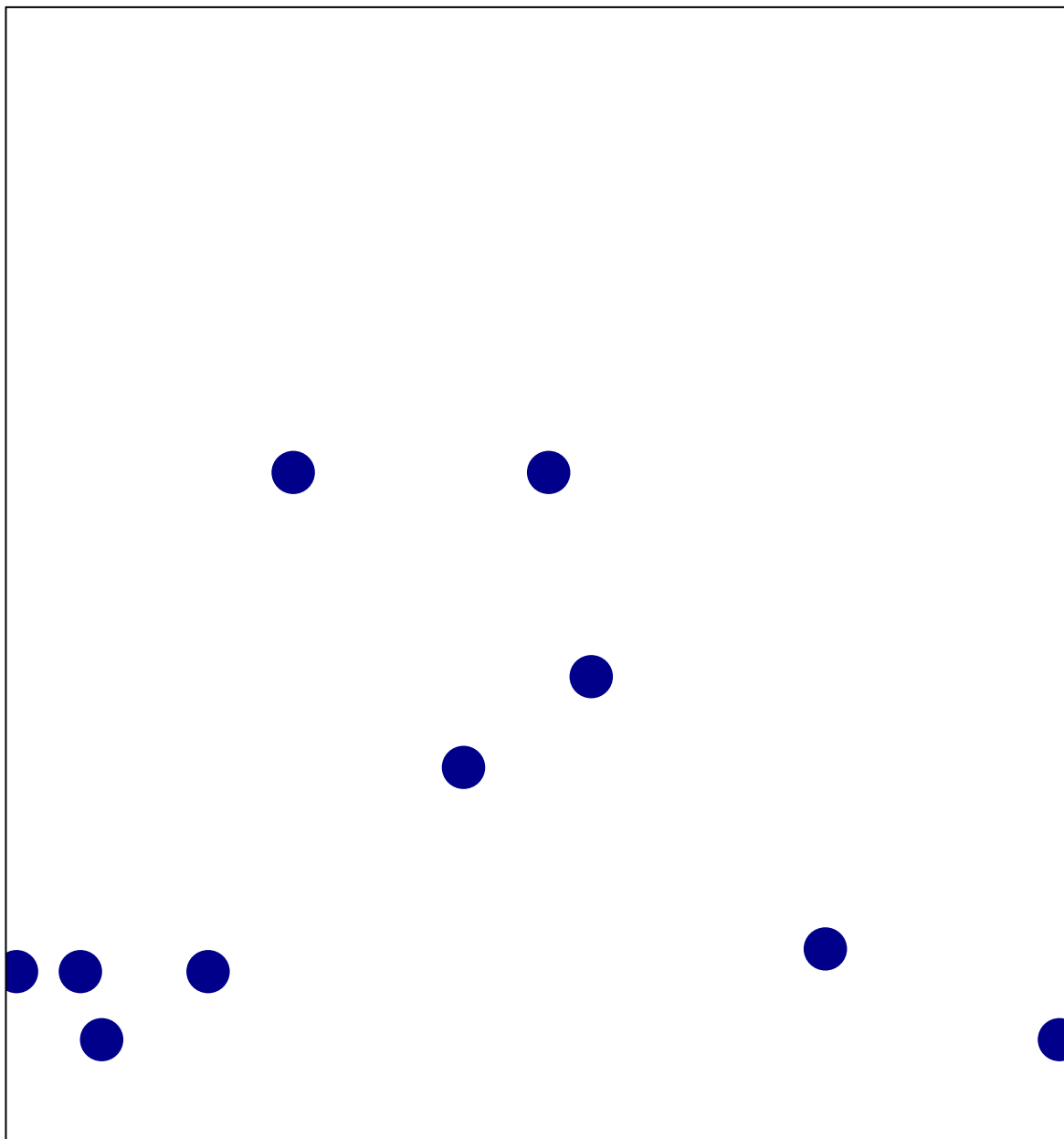
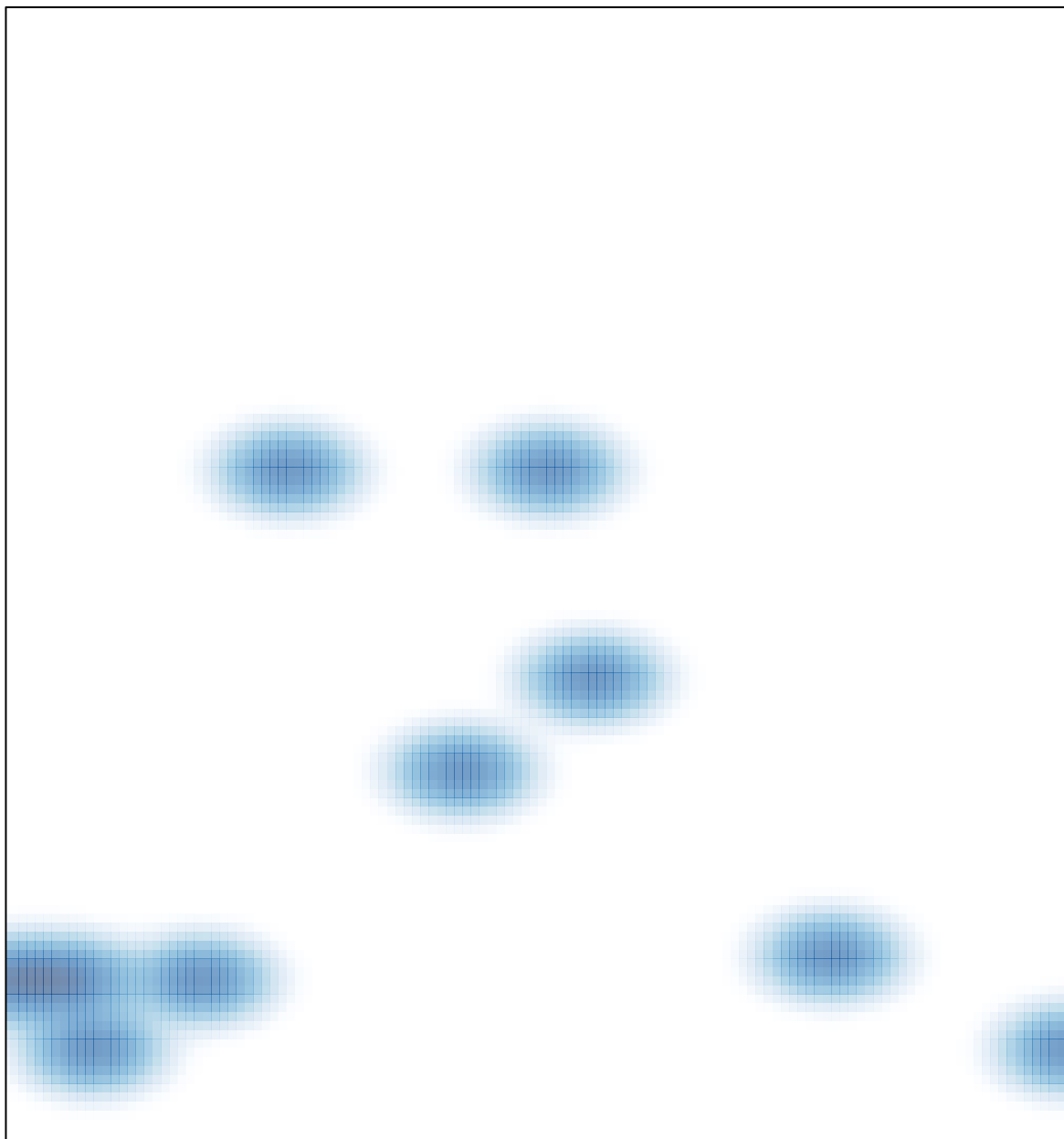


# REACTOME\_HDL\_MEDIATED\_LIPID\_TRANSPORT



# features = 10  
chi-square p = 0.82

# REACTOME\_HDL\_MEDIATED\_LIPID\_TRANSPORT



# features = 10 , max = 1