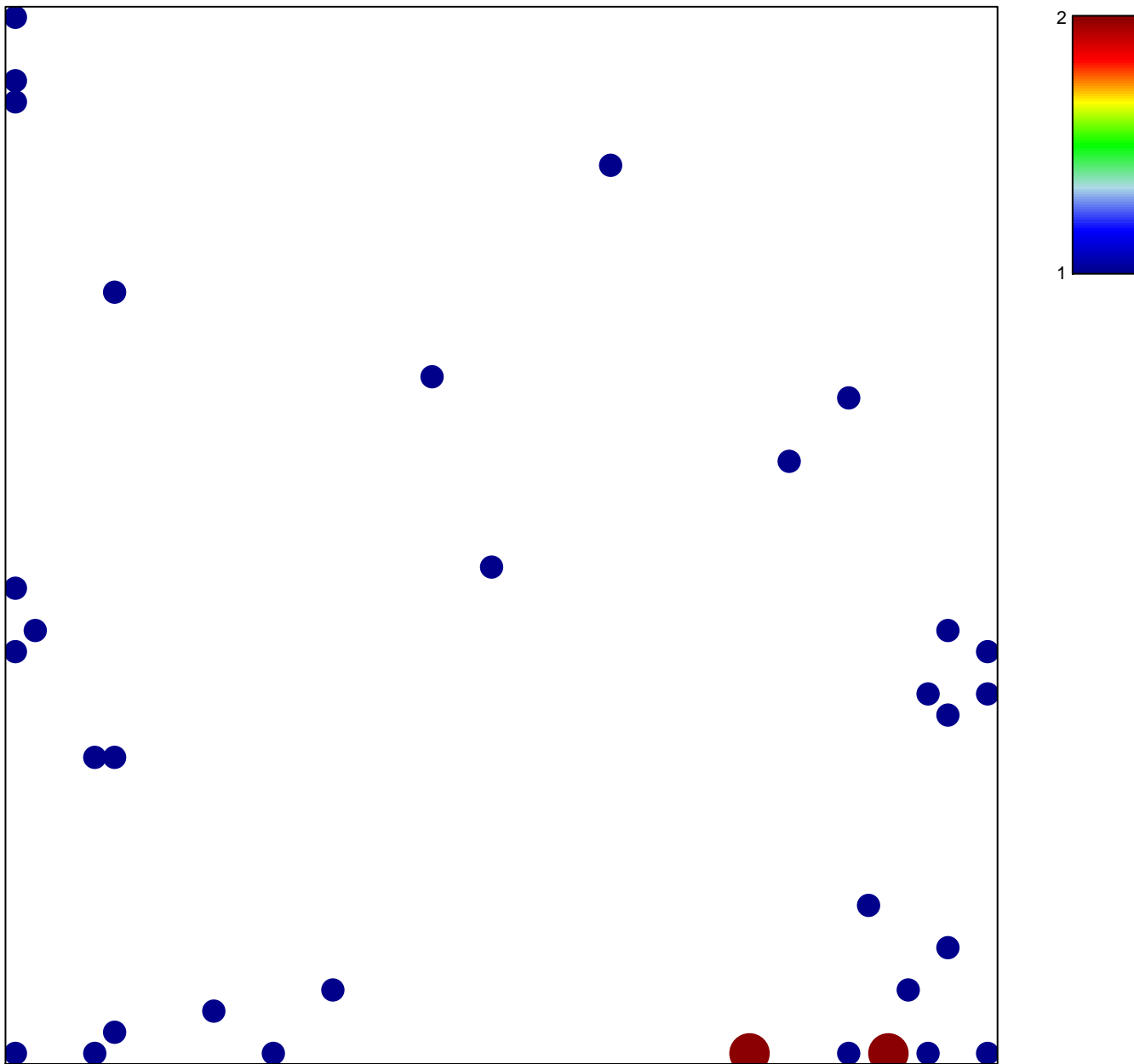
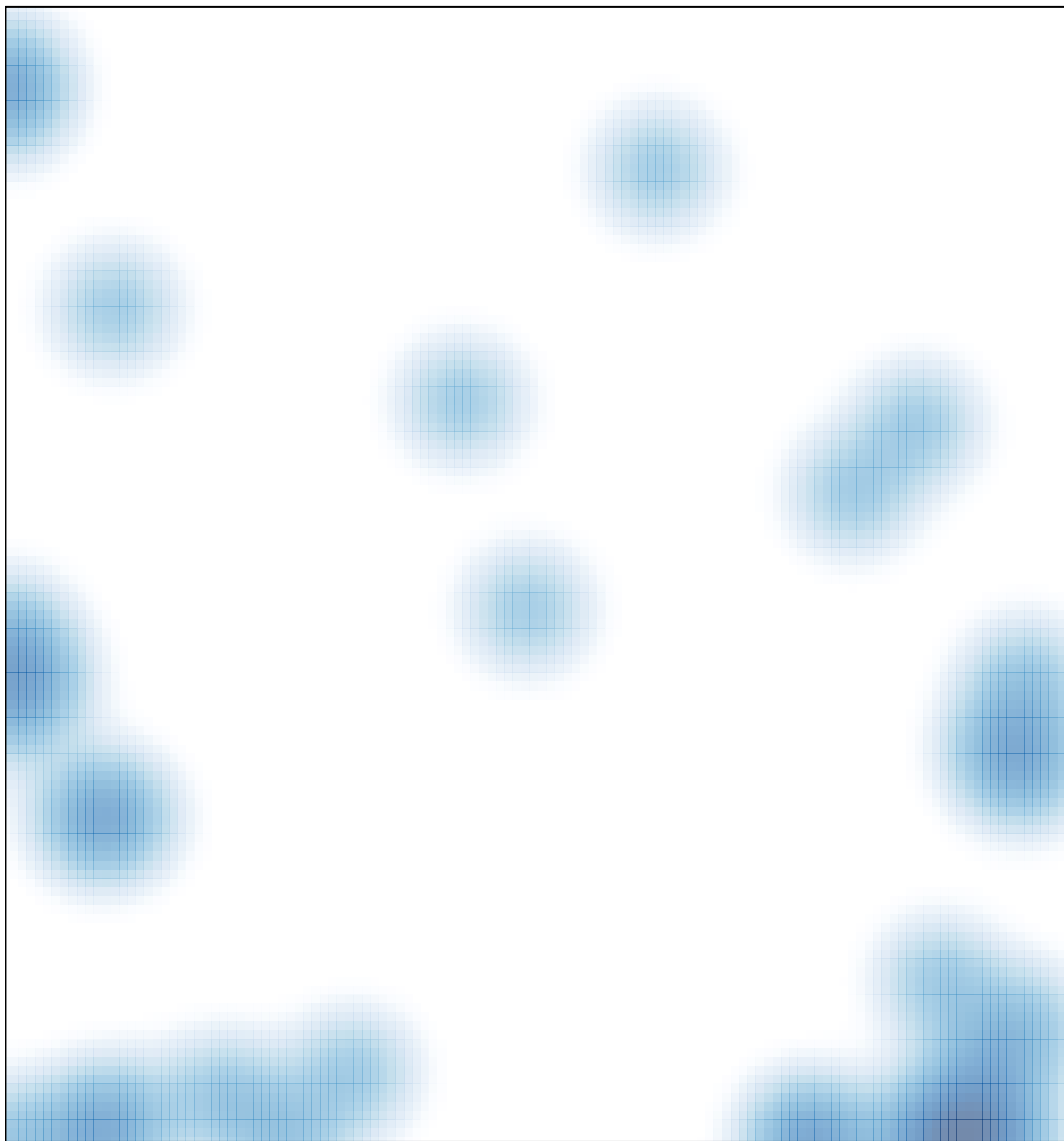


# REACTOME\_TRANSCRIPTIONAL\_ACTIVITY\_OF\_SMAD2\_SMAD3\_SMAD4\_HETEROTRIMER



# features = 35  
chi-square p = 0.75

# SCRIPTONAL\_ACTIVITY\_OF\_SMAD2\_SMAD3\_SMAI



# features = 35 , max = 2