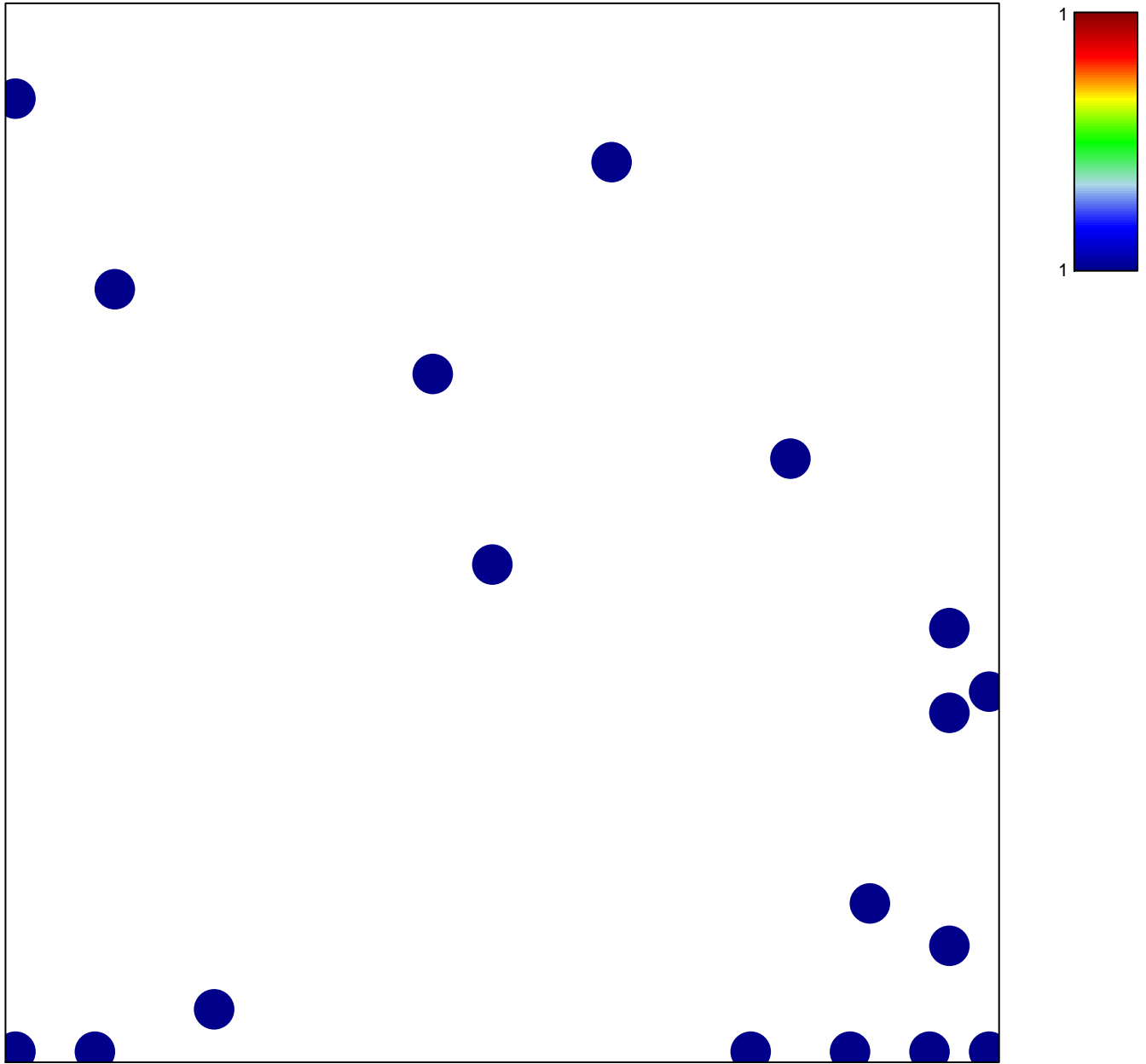
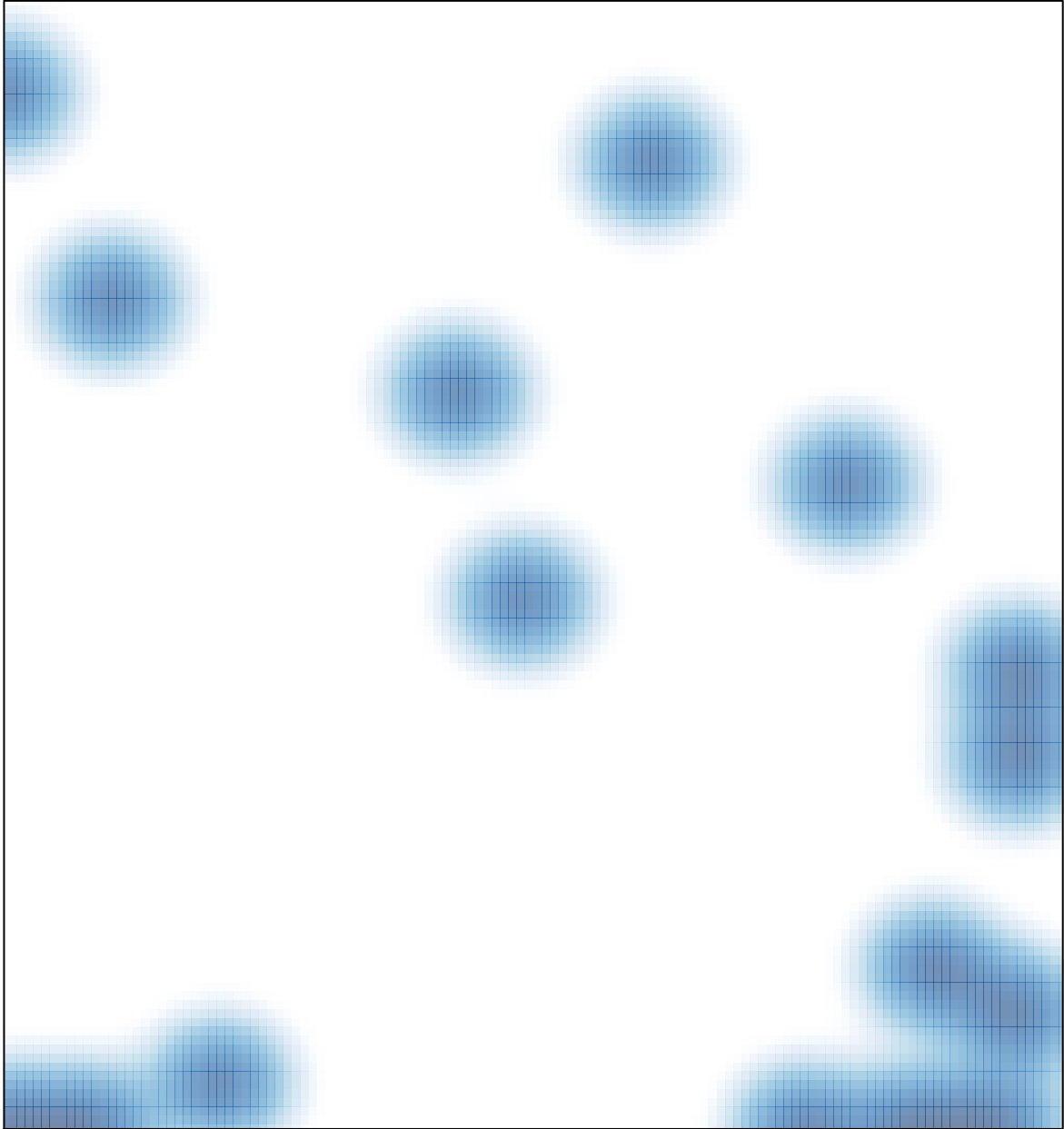


# REACTOME\_DOWNREGULATION\_OF\_SMAD2\_3\_SMAD4\_TRANSCRIPTIONAL\_ACTIVITY



# features = 18  
chi-square p = 0.79

# UNREGULATION\_OF\_SMAD2\_3\_SMAD4\_TRANSCRIP



# features = 18 , max = 1