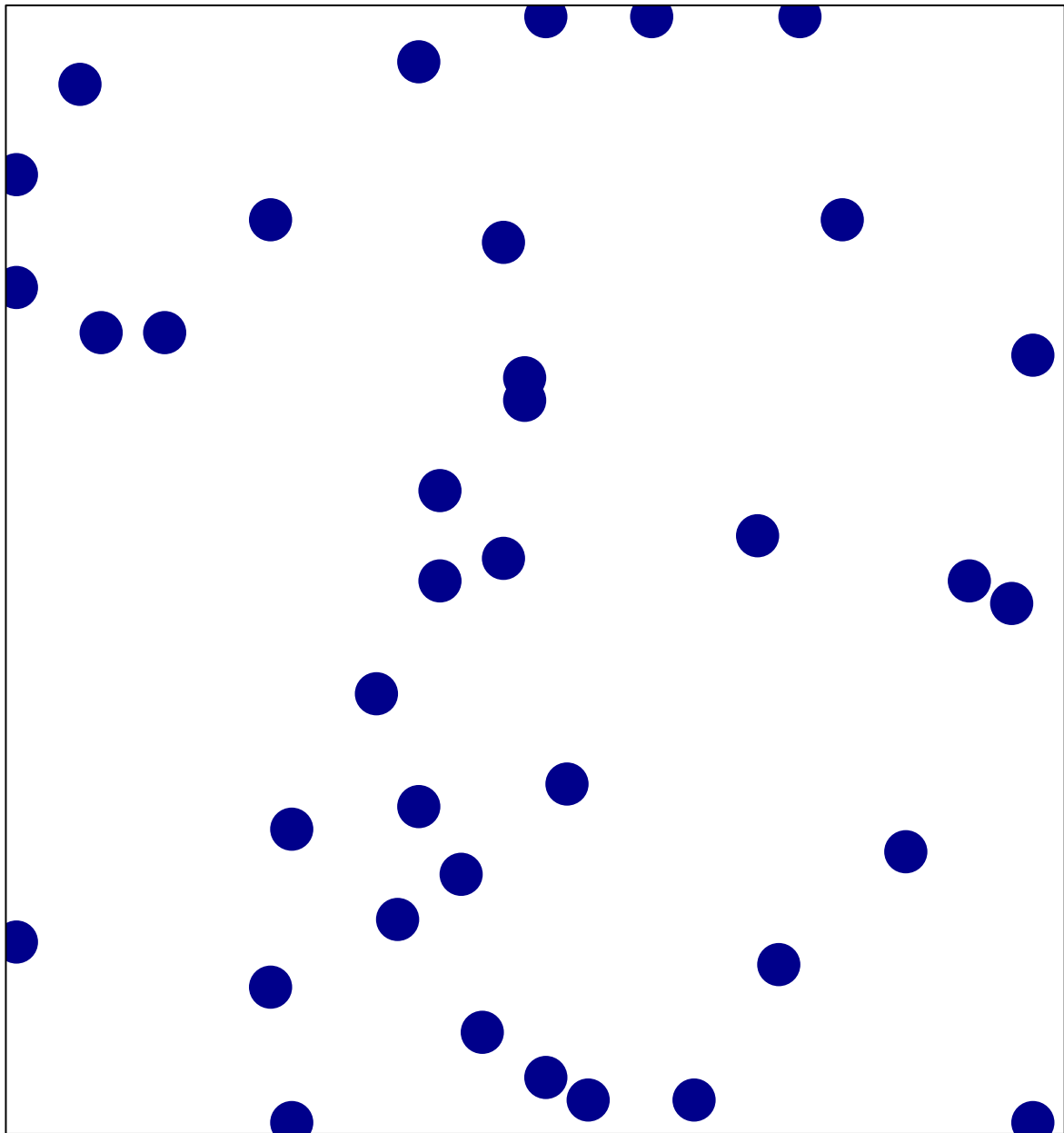


# NOUSHMEHR\_GBM\_SILENCED\_BY\_METHYLATION



# features = 37  
chi-square p = 0.86

# NOUSHMEHR\_GBM\_SILENCED\_BY\_METHYLATION



# features = 37 , max = 1