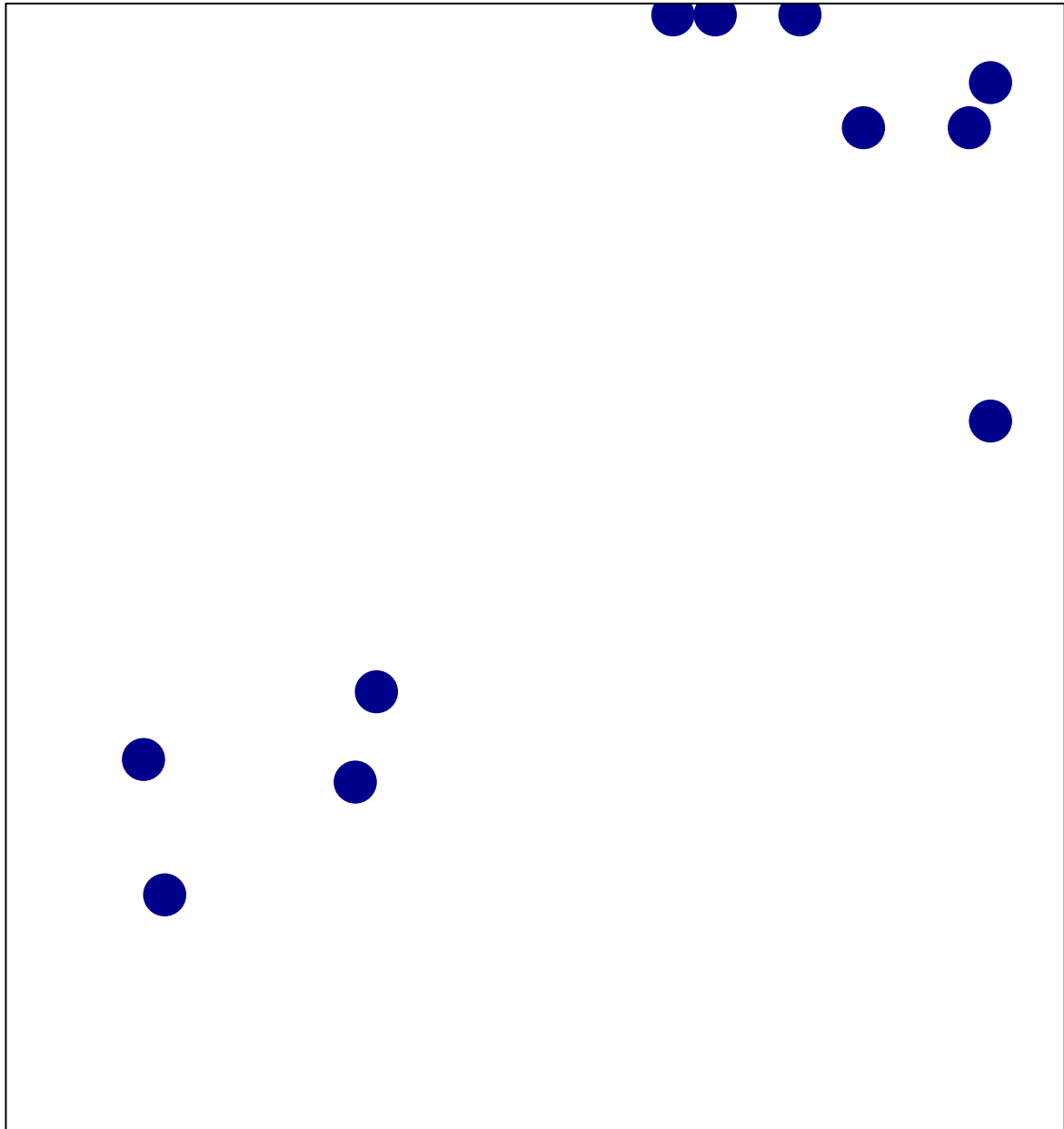


# REACTOME\_MRNA\_DECAY\_BY\_5\_TO\_3\_EXORIBONUCLEASE



# features = 11  
chi-square p = 0.82

# CTOME\_MRNA\_DECAY\_BY\_5\_TO\_3\_EXORIBONUCL



# features = 11 , max = 1