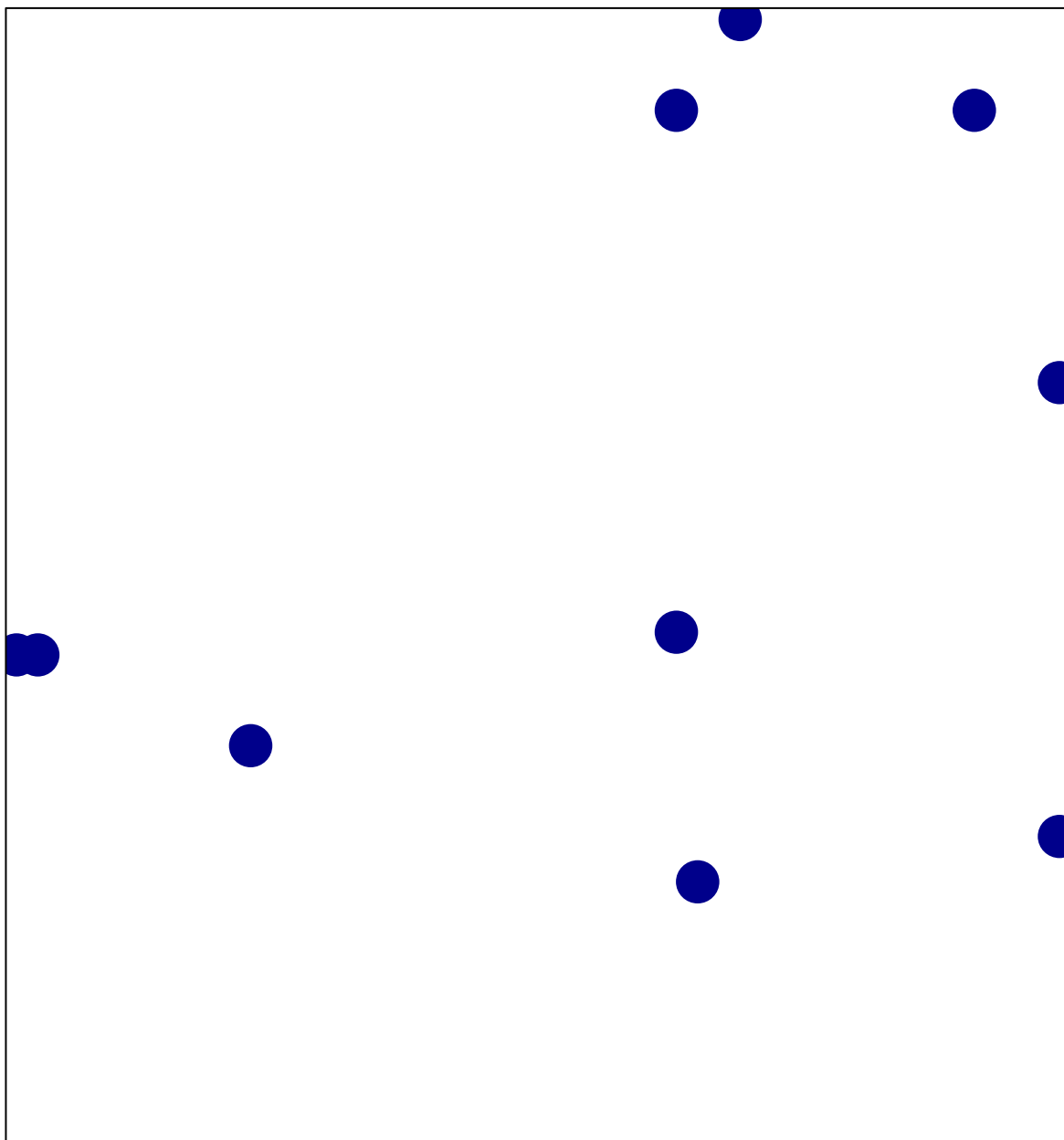


# REACTOME\_MITOCHONDRIAL\_FATTY\_ACID\_BETA\_OXIDATION



# features = 10  
chi-square p = 0.83

# CTOME\_MITOCHONDRIAL\_FATTY\_ACID\_BETA\_OXID



# features = 10 , max = 1