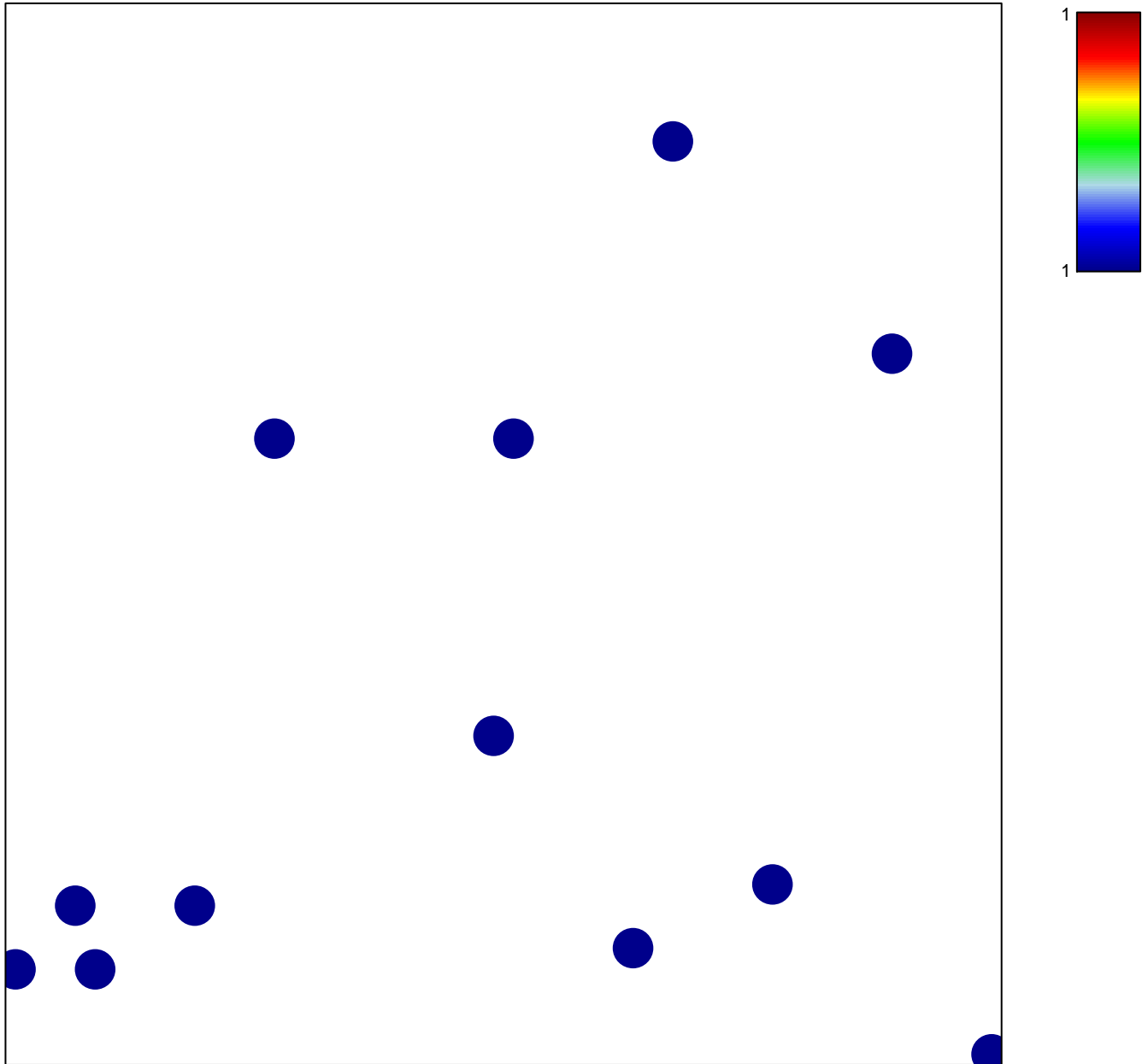


# reverse cholesterol transport



# features = 12  
chi-square p = 0.84

# reverse cholesterol transport



# features = 12 , max = 1