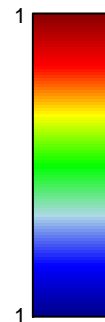
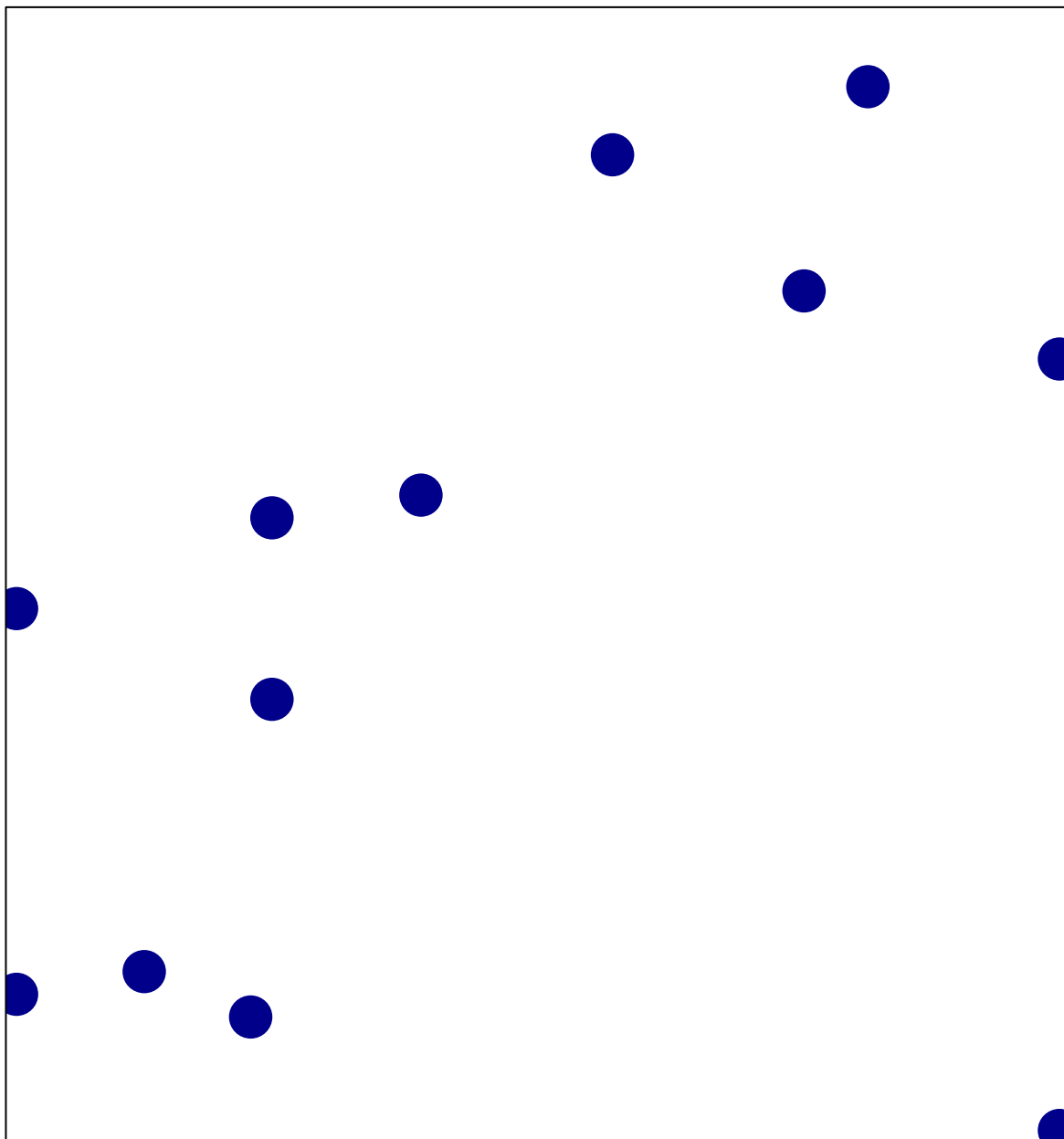


# low-density lipoprotein particle receptor binding



# features = 12  
chi-square p = 0.84

# low-density lipoprotein particle receptor binding



# features = 12 , max = 1