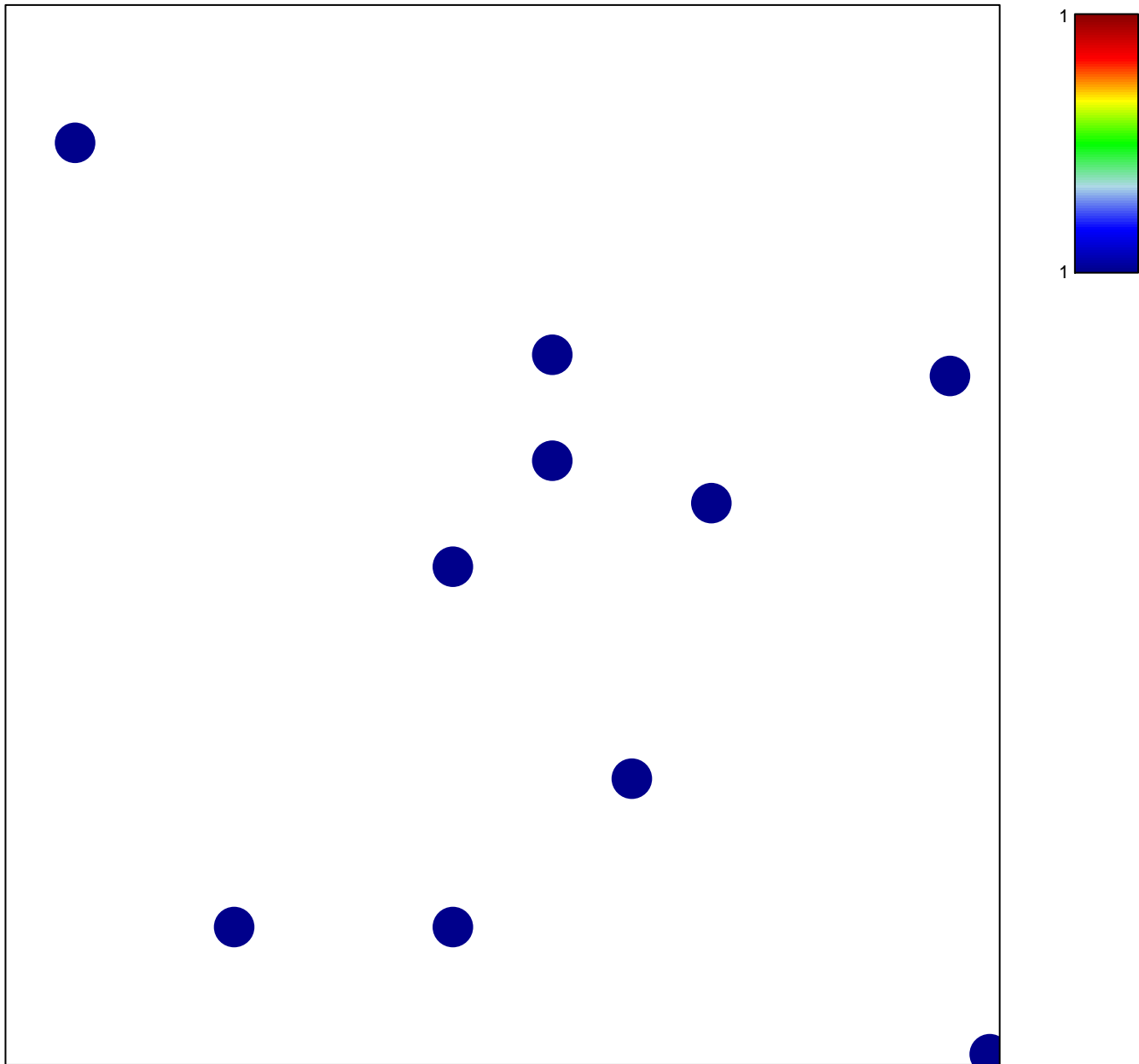


# tropomyosin binding



# features = 10  
chi-square p = 0.83

# tropomyosin binding



# features = 10 , max = 1