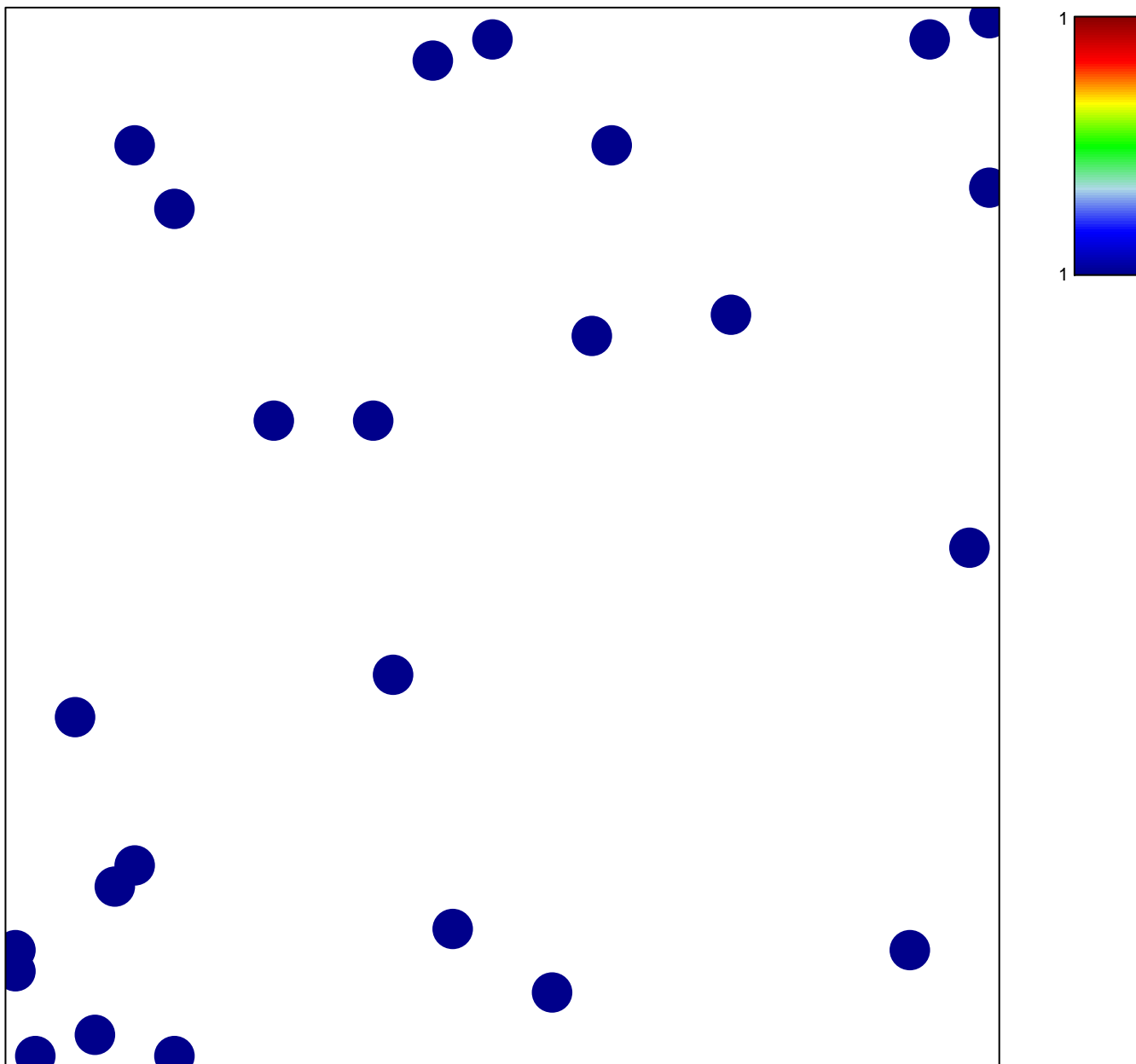
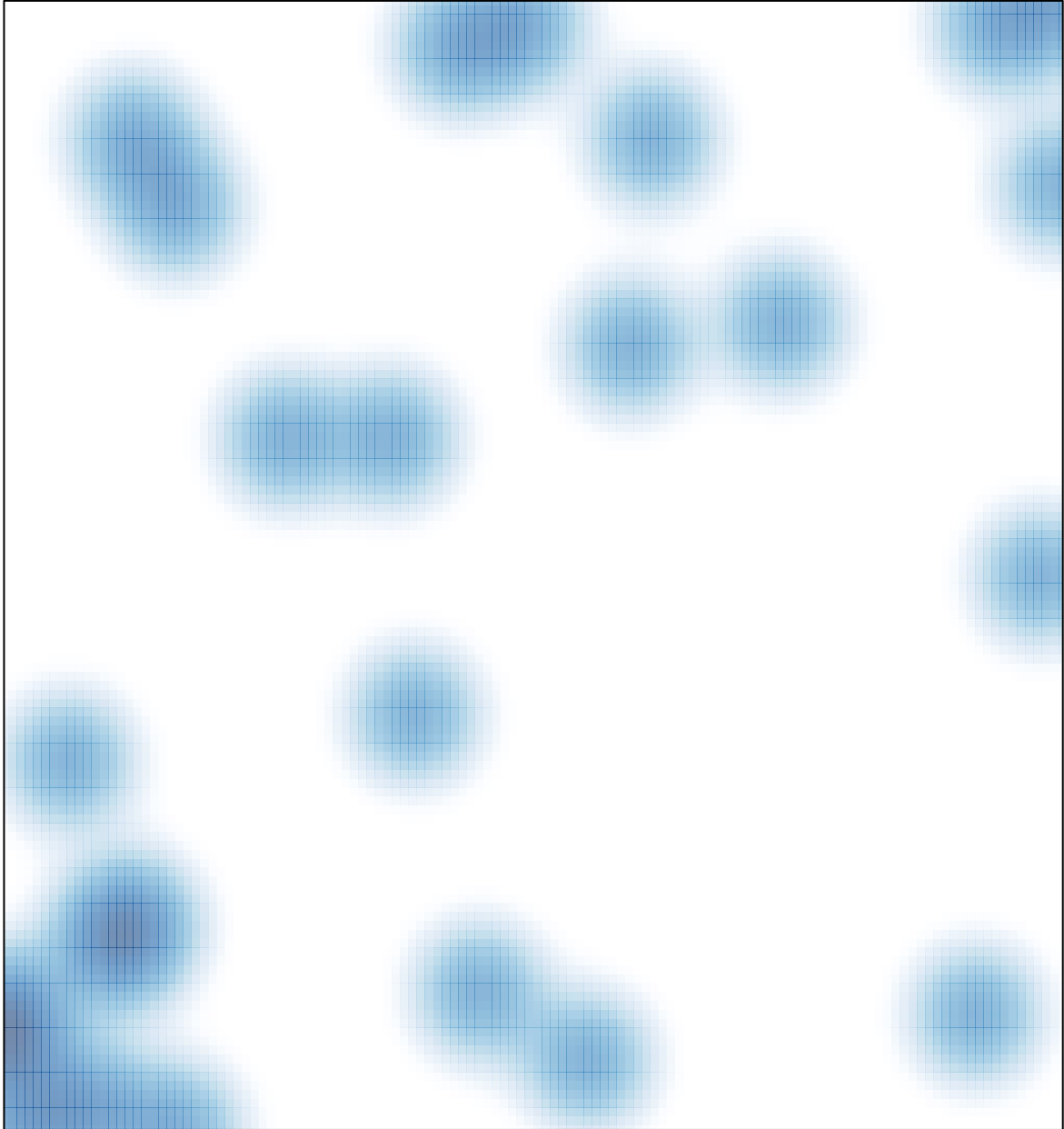


# protein localization to nucleus



# features = 25  
chi-square p = 0.85

# protein localization to nucleus



# features = 25 , max = 1