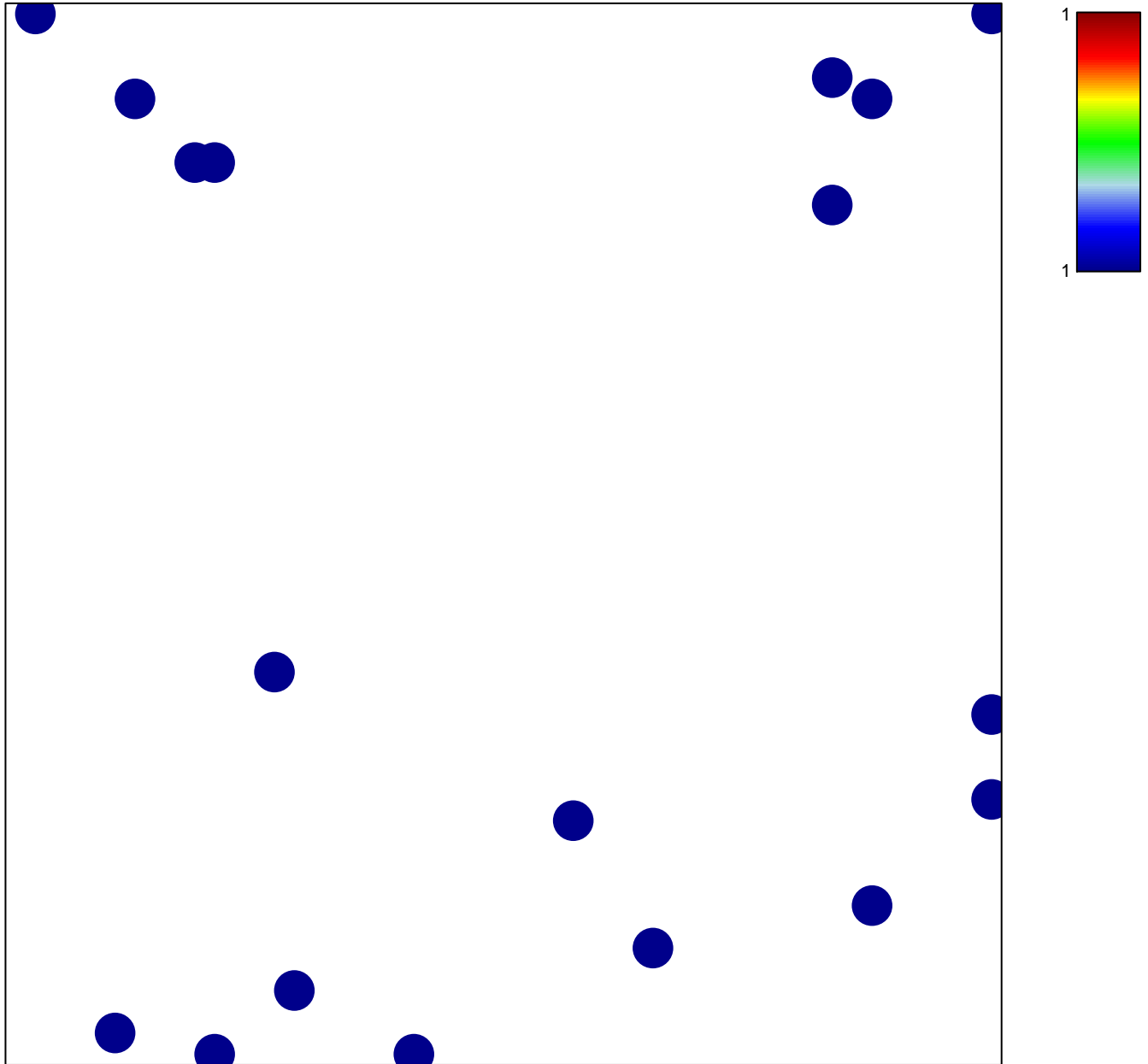
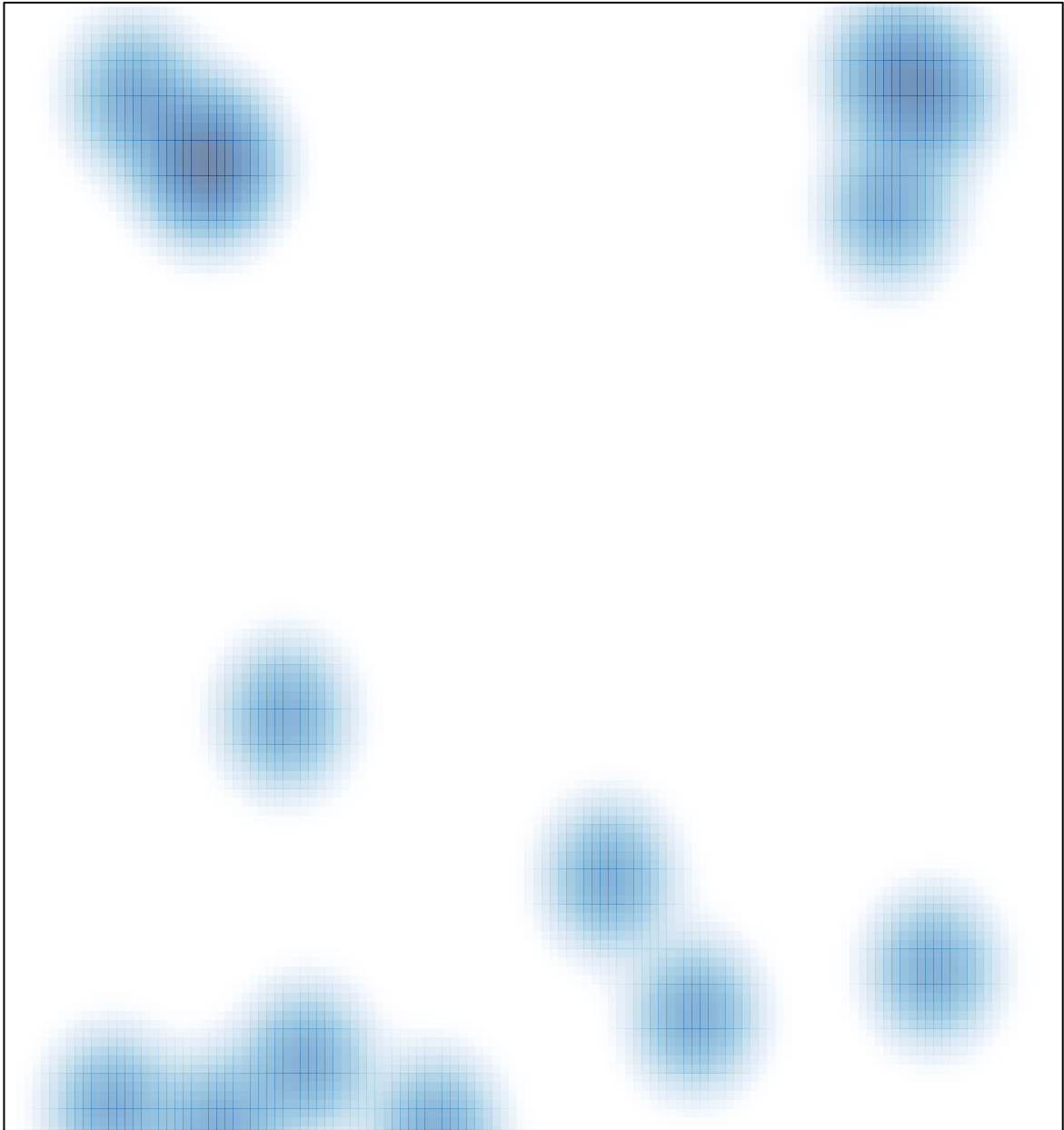


# histone lysine methylation



# features = 18  
chi-square p = 0.83

# histone lysine methylation



# features = 18 , max = 1