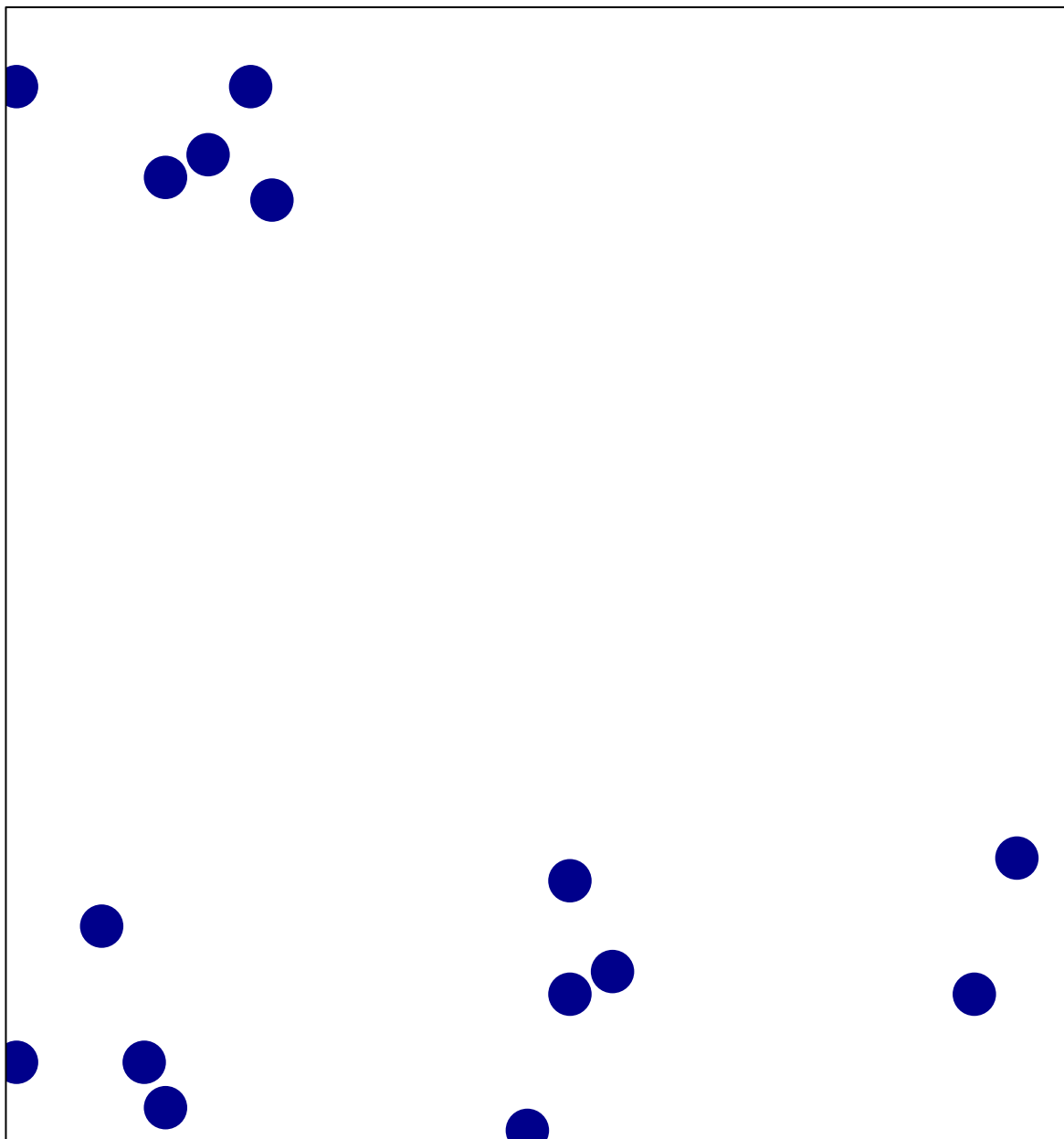
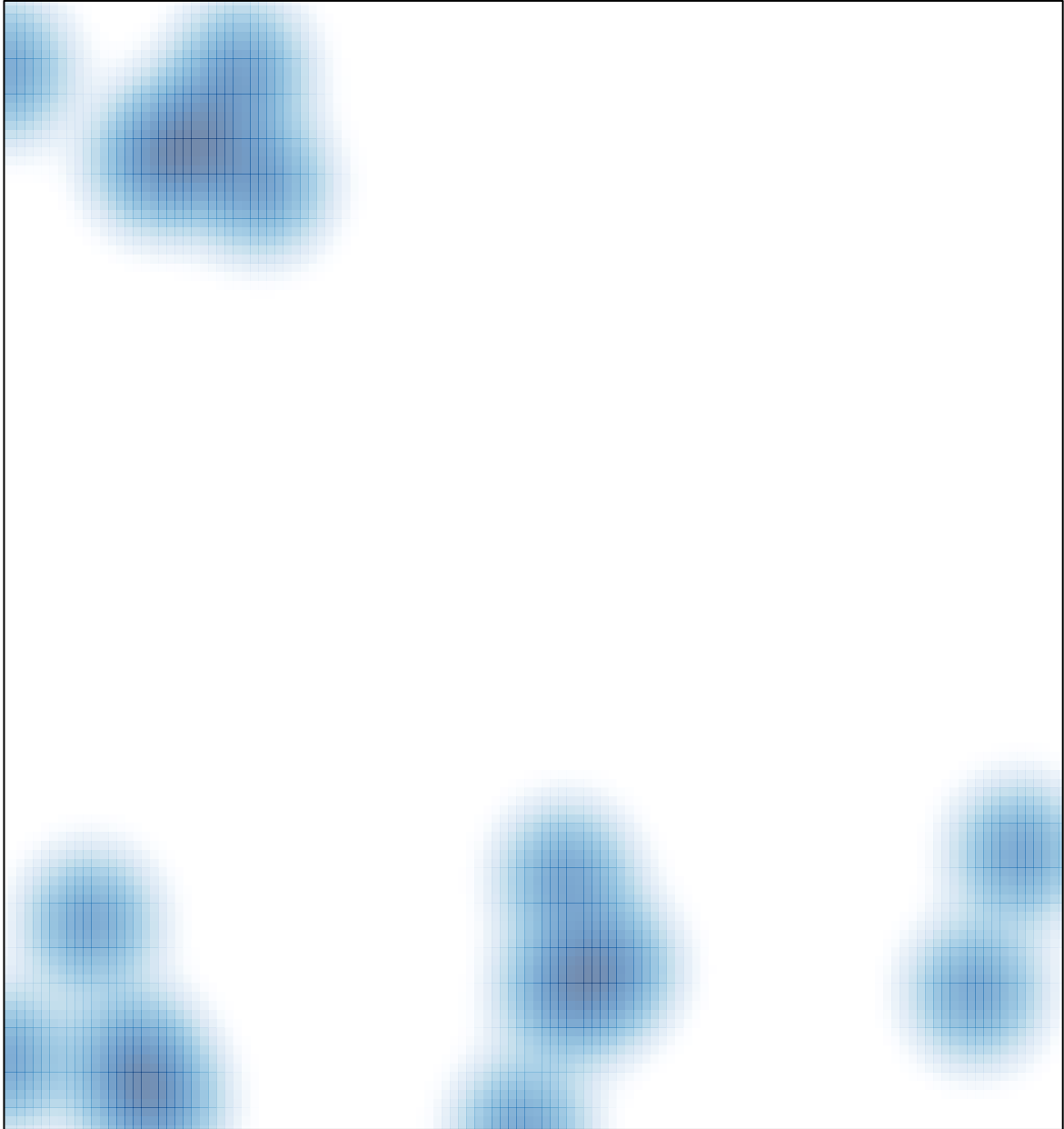


# lysine-acetylated histone binding



# features = 15  
chi-square p = 0.83

# lysine-acetylated histone binding



# features = 15 , max = 1