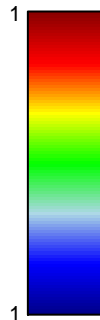
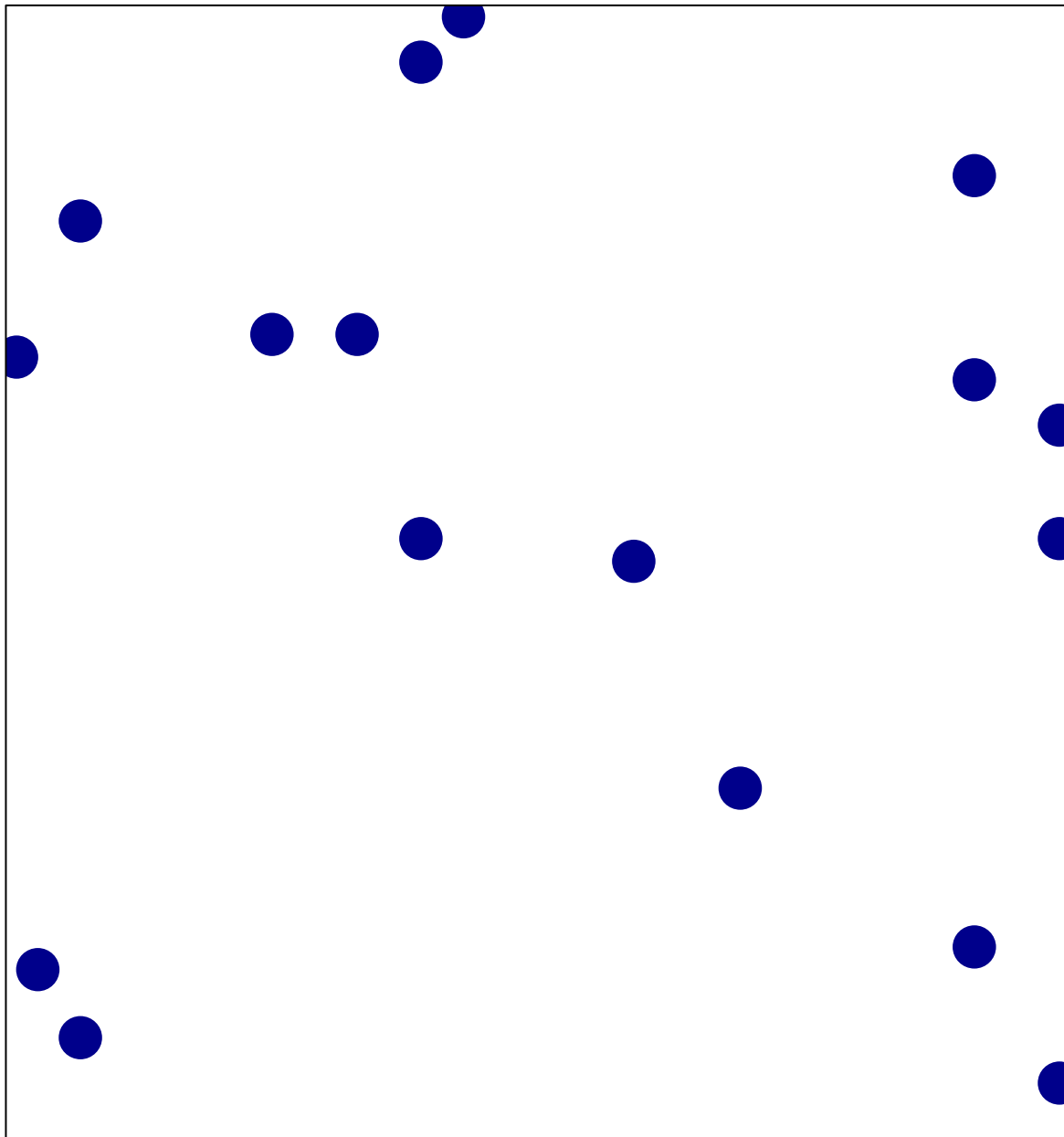
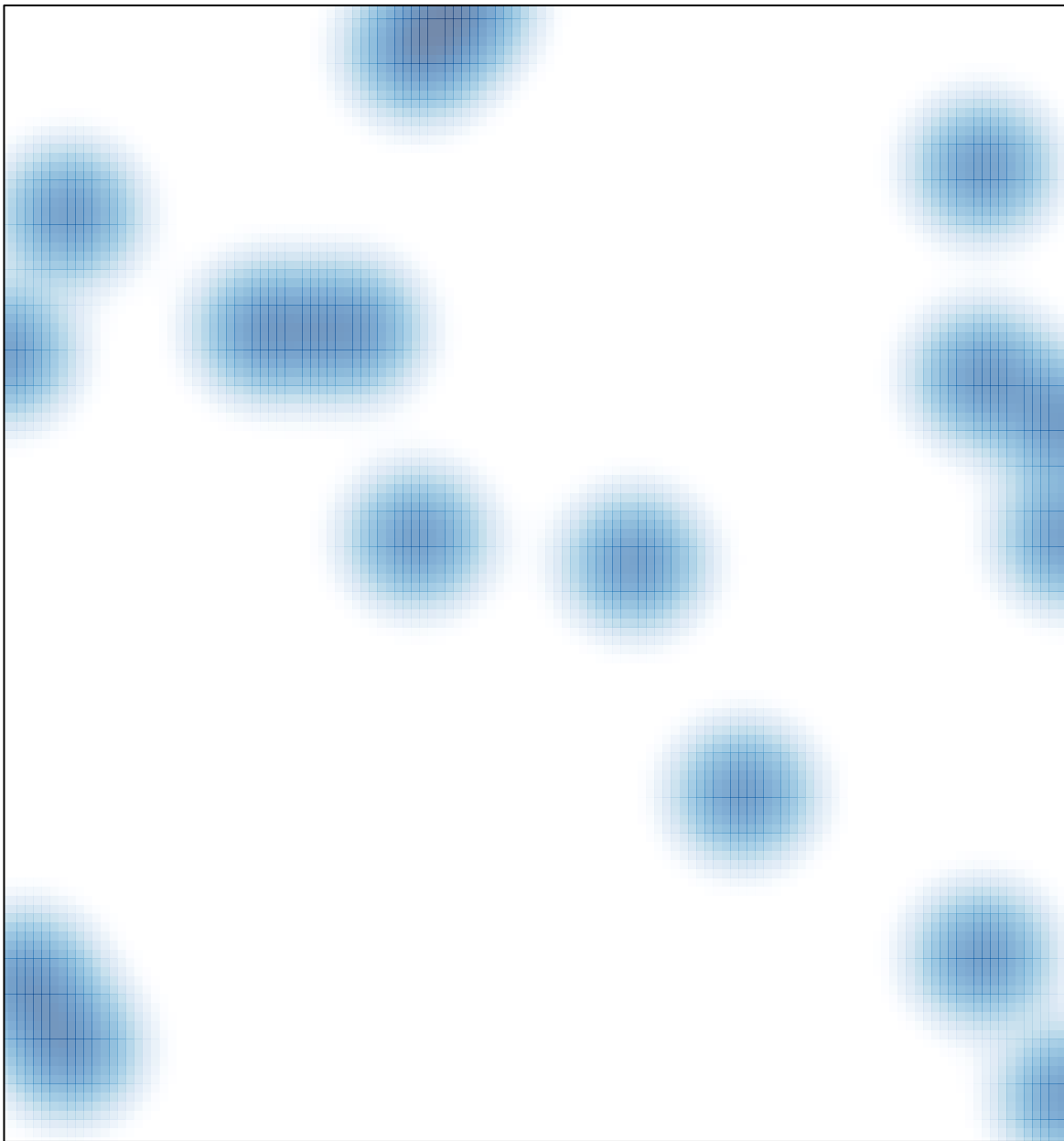


# histone acetyltransferase binding



# features = 17  
chi-square p = 0.82

# histone acetyltransferase binding



# features = 17 , max = 1