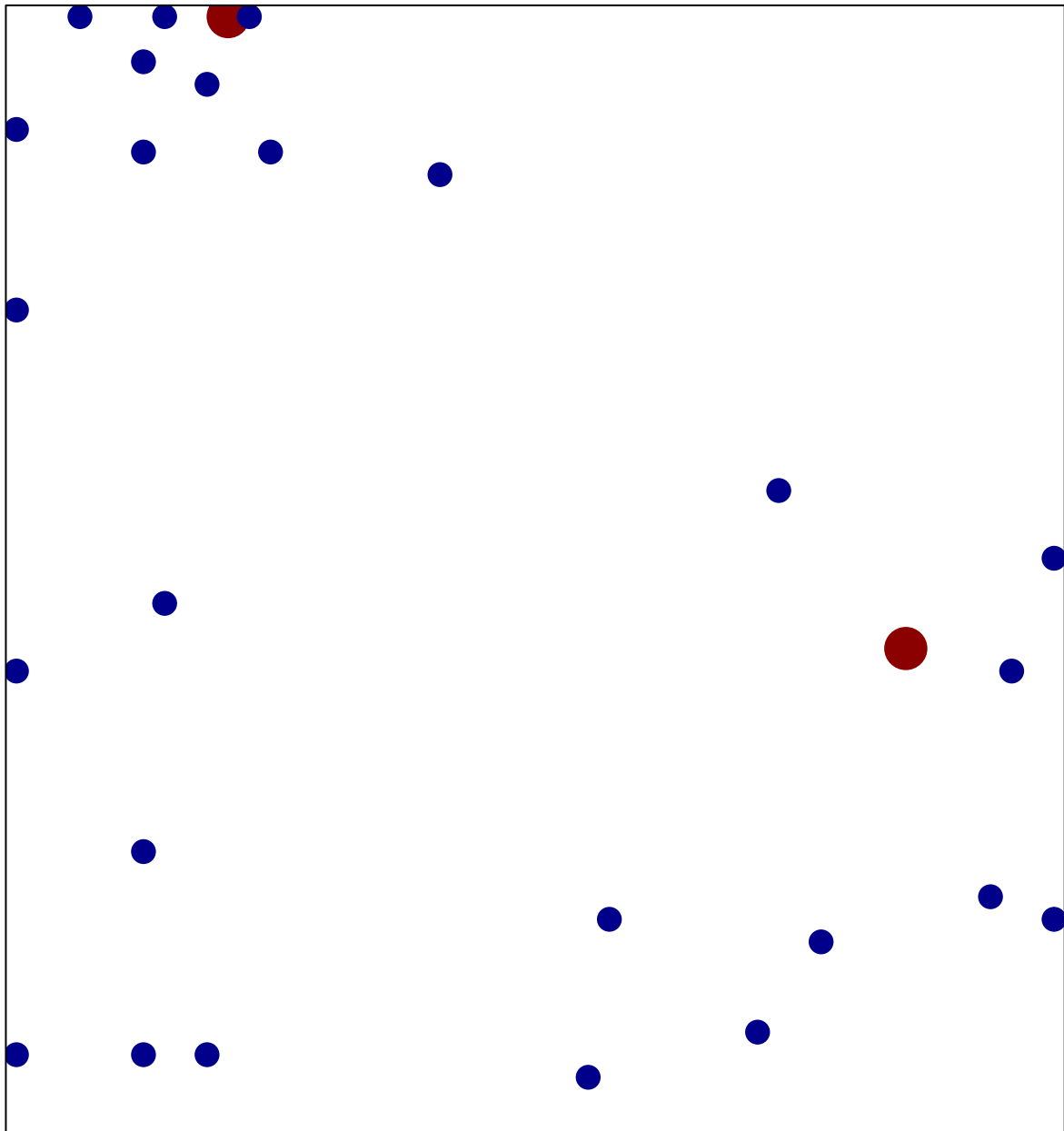
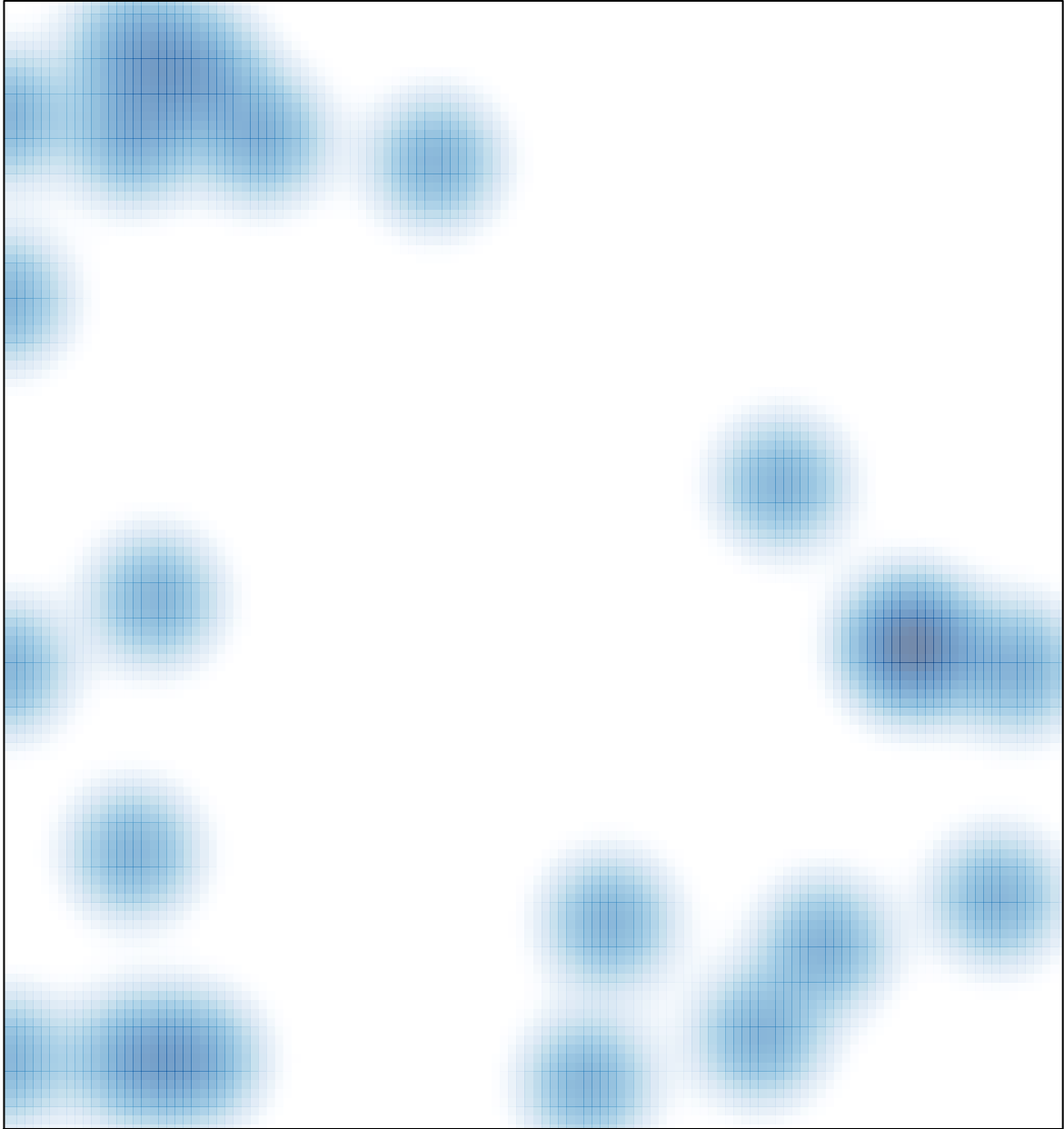


# nucleosomal DNA binding



# features = 29  
chi-square p = 0.8

# nucleosomal DNA binding



# features = 29 , max = 2