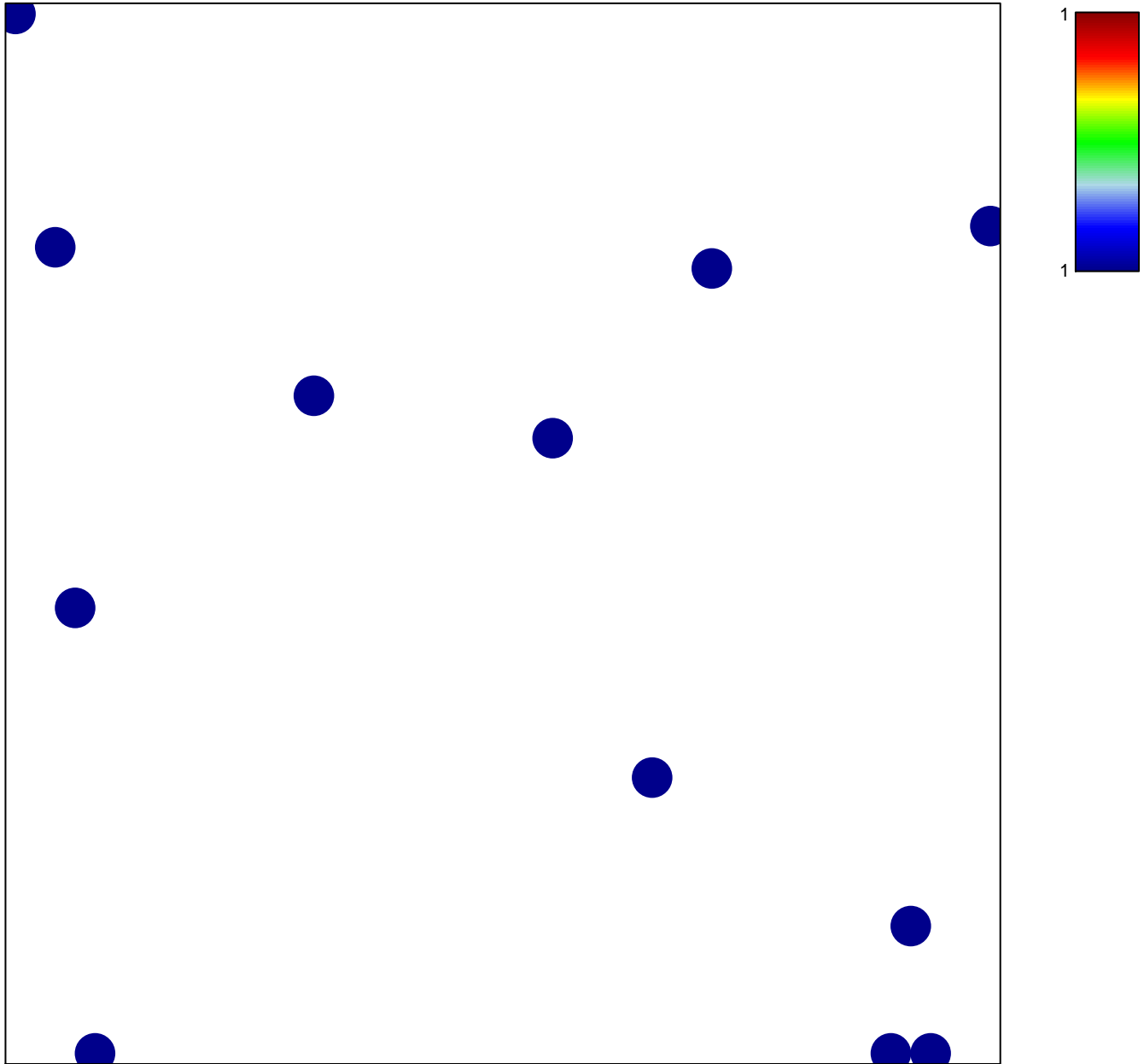
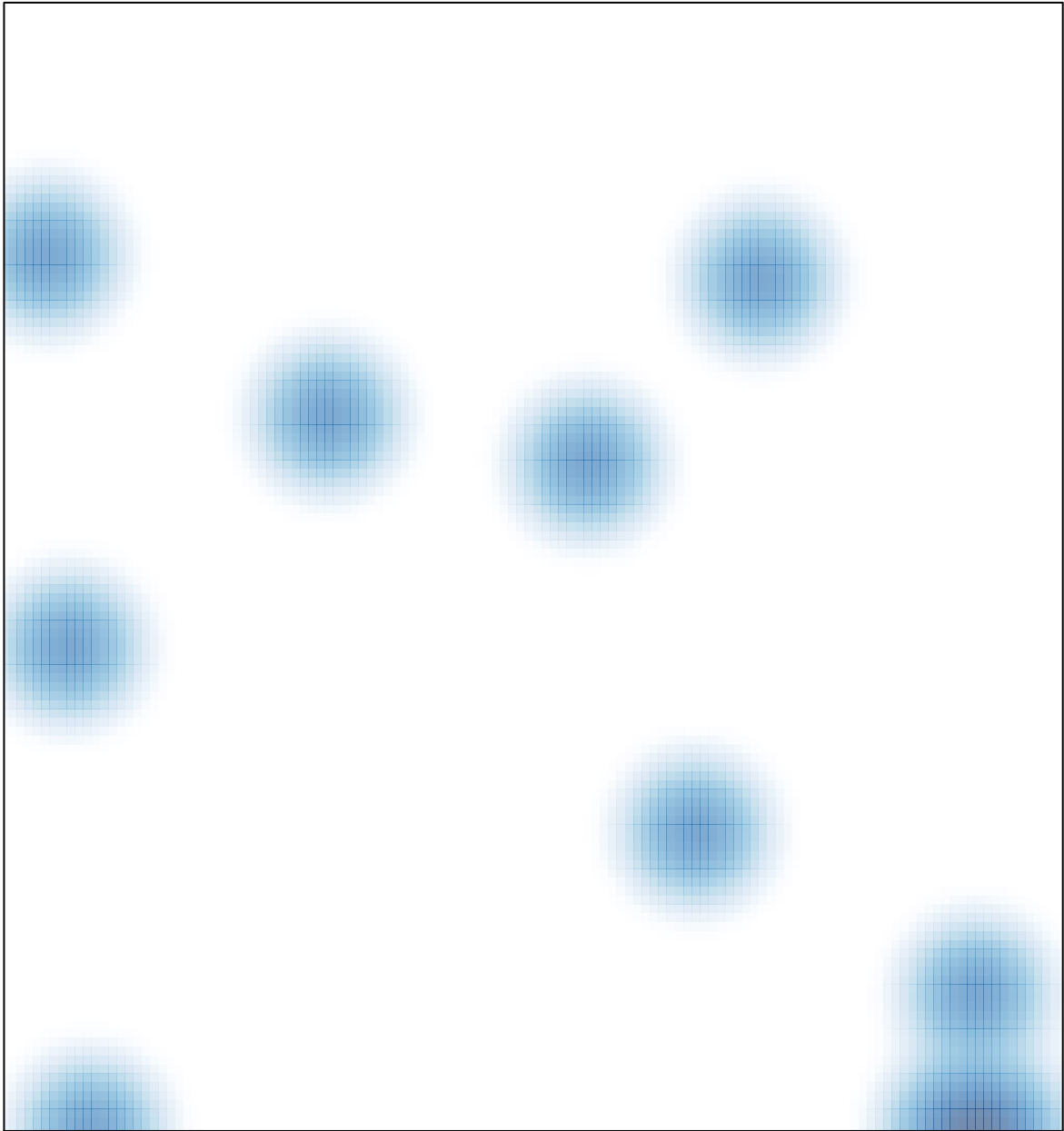


# core promoter proximal region sequence-specific DNA binding



# features = 12  
chi-square p = 0.84

# promoter proximal region sequence-specific DNA b



# features = 12 , max = 1