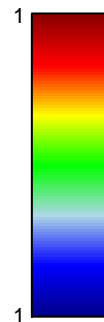
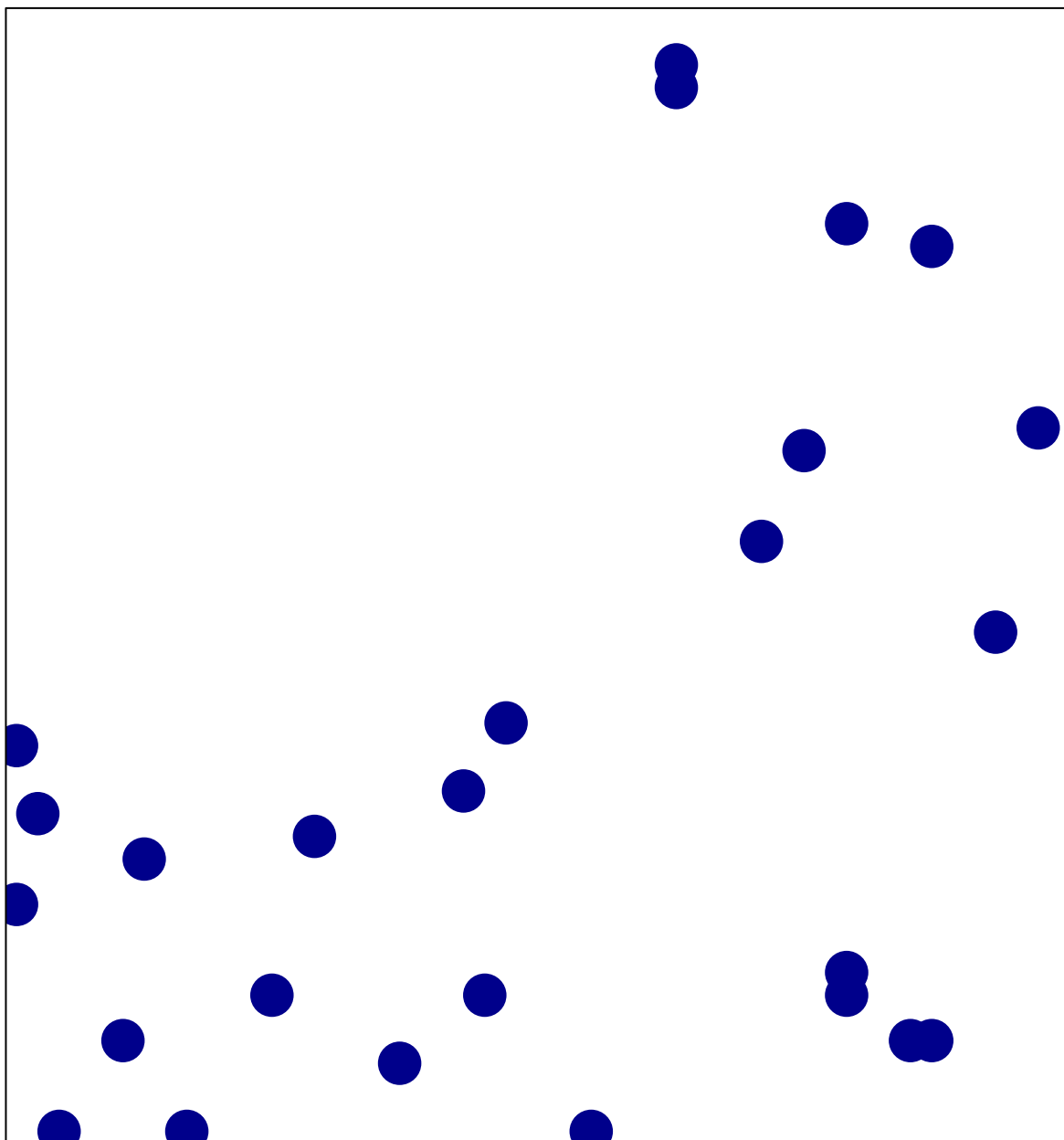
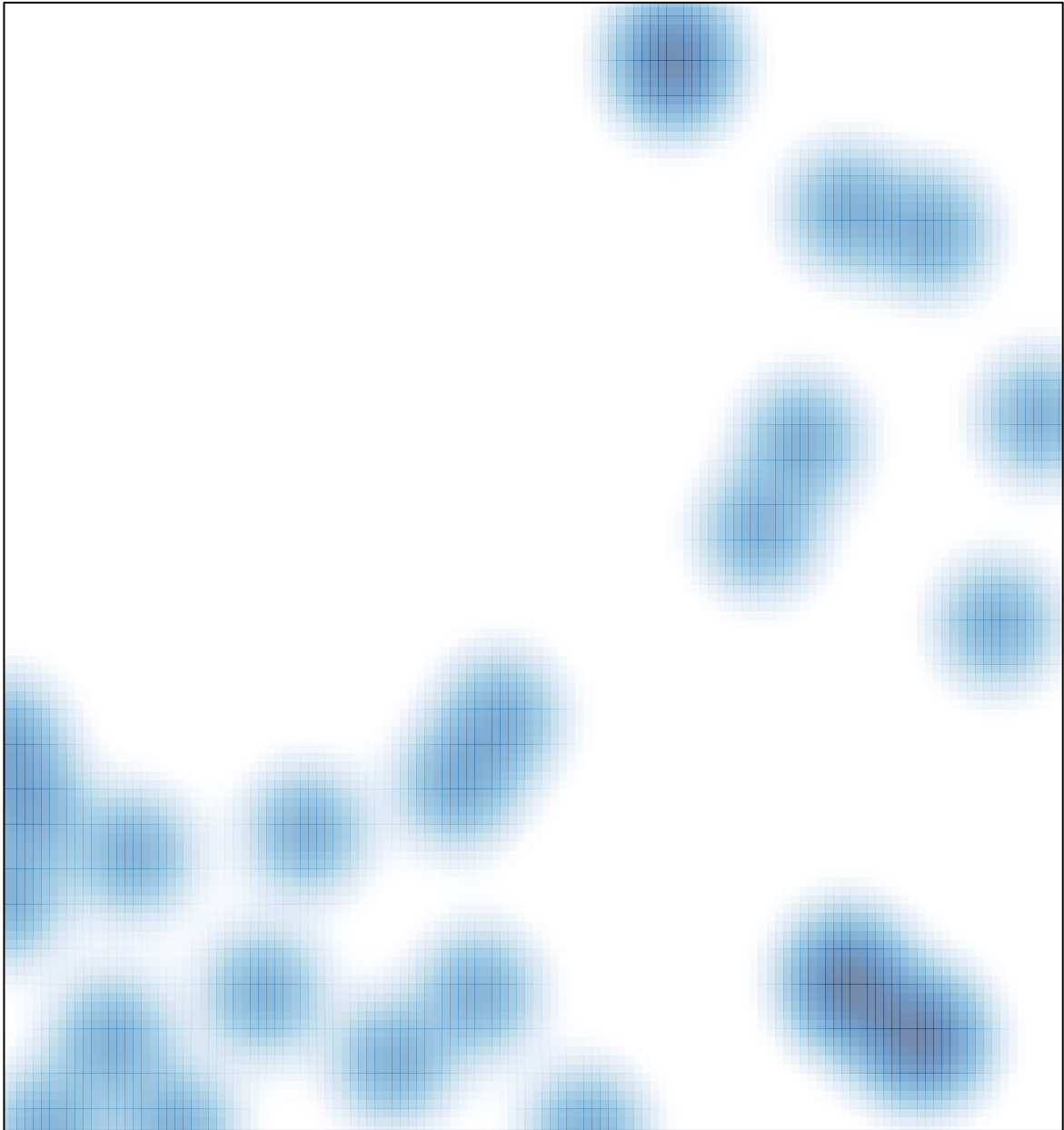


# peptide antigen binding



# features = 26  
chi-square p = 0.84

# peptide antigen binding



# features = 26 , max = 1