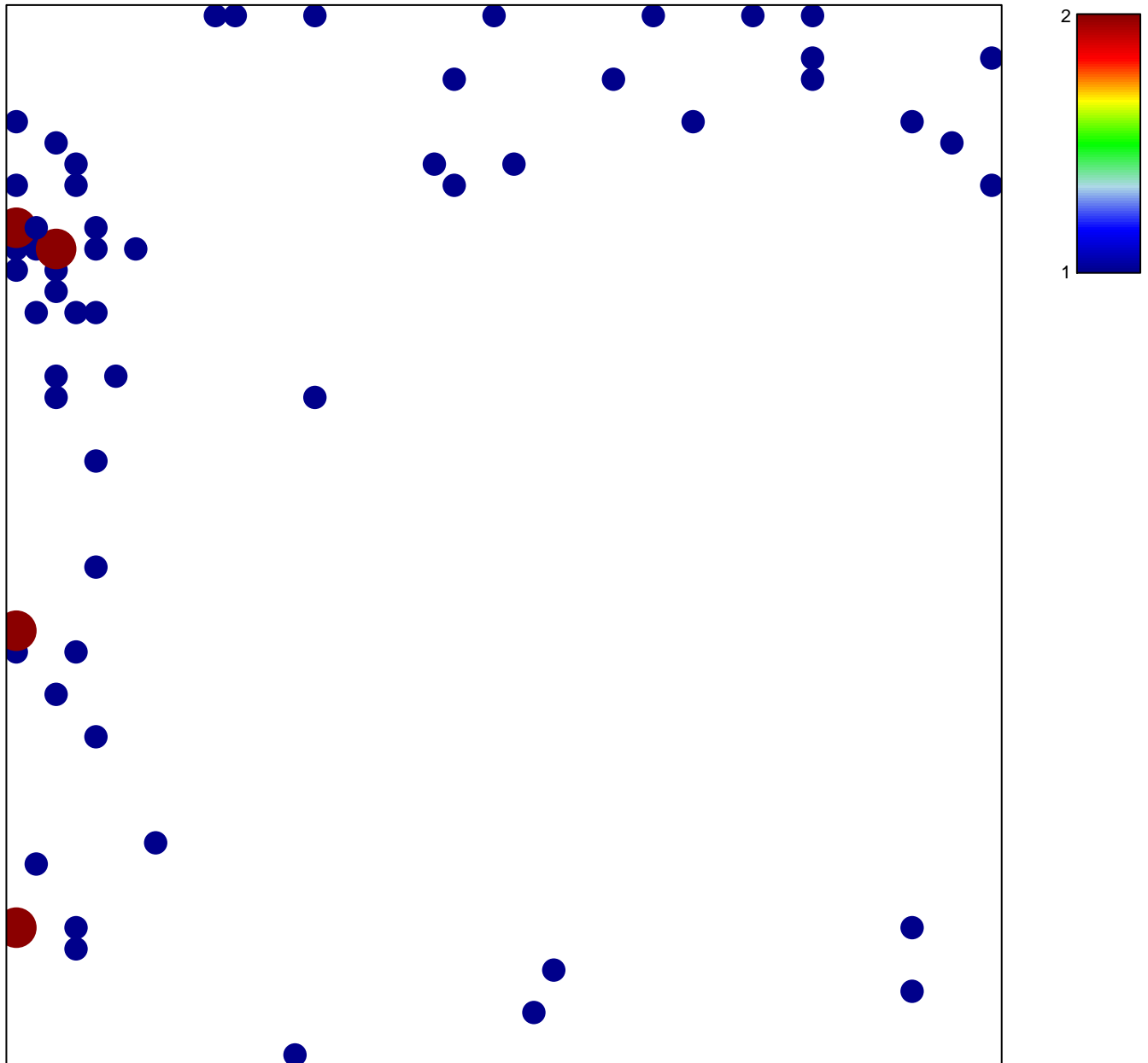
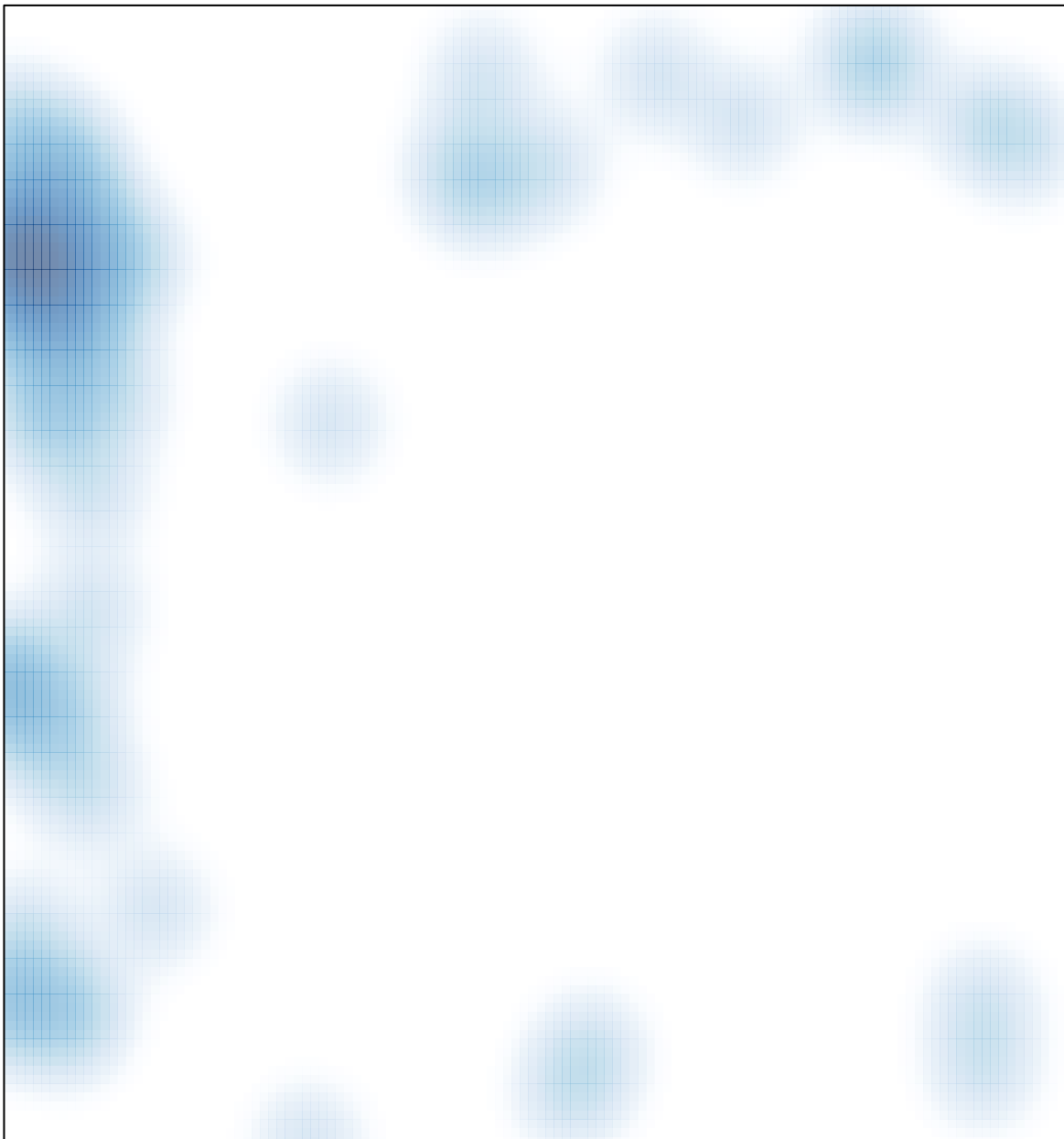


# Stuehler\_Proteins\_up\_in\_STS



# features = 63  
chi-square p = 0.73

# Stuehler\_Proteins\_up\_in\_STS



# features = 63 , max = 2