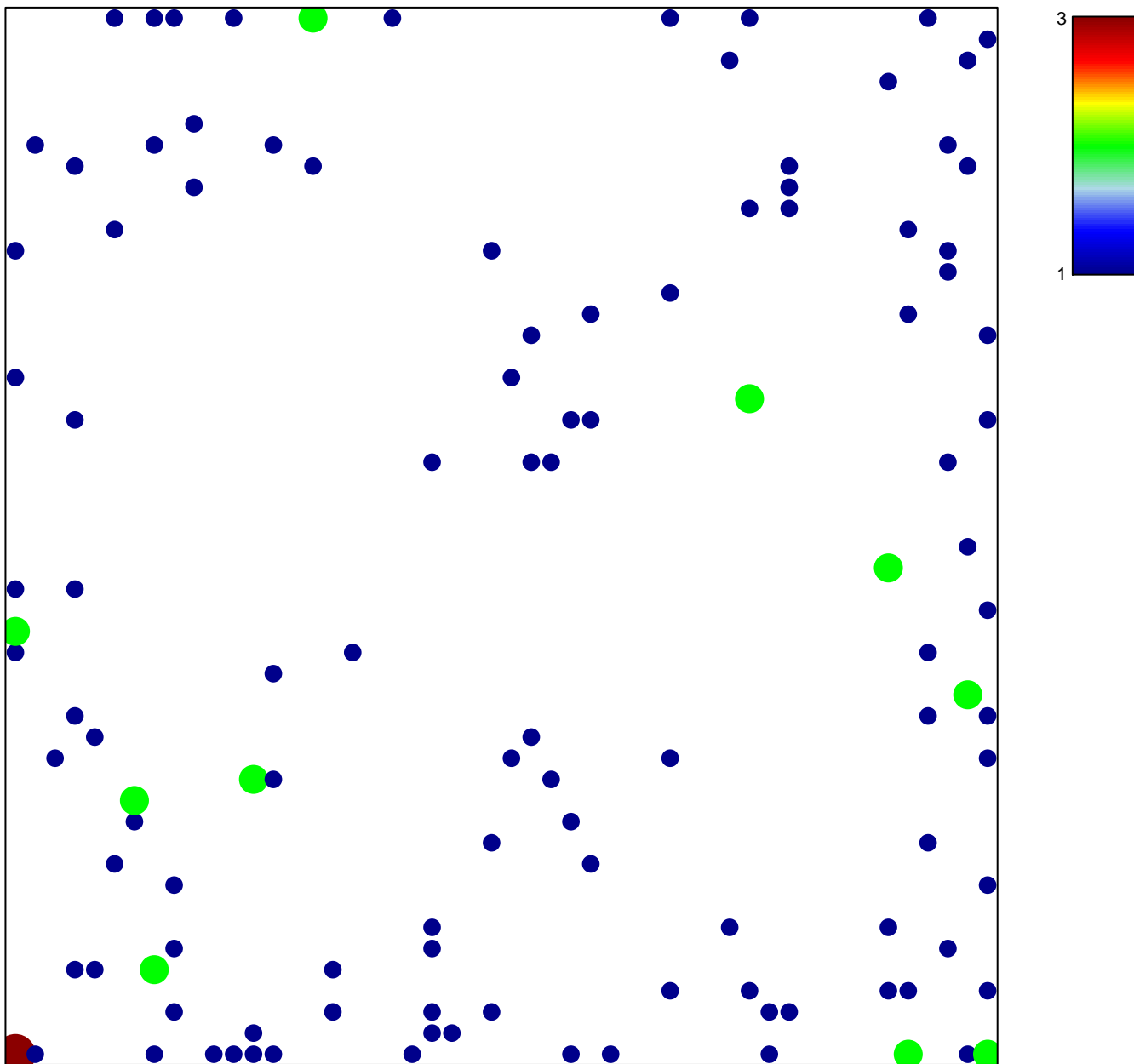
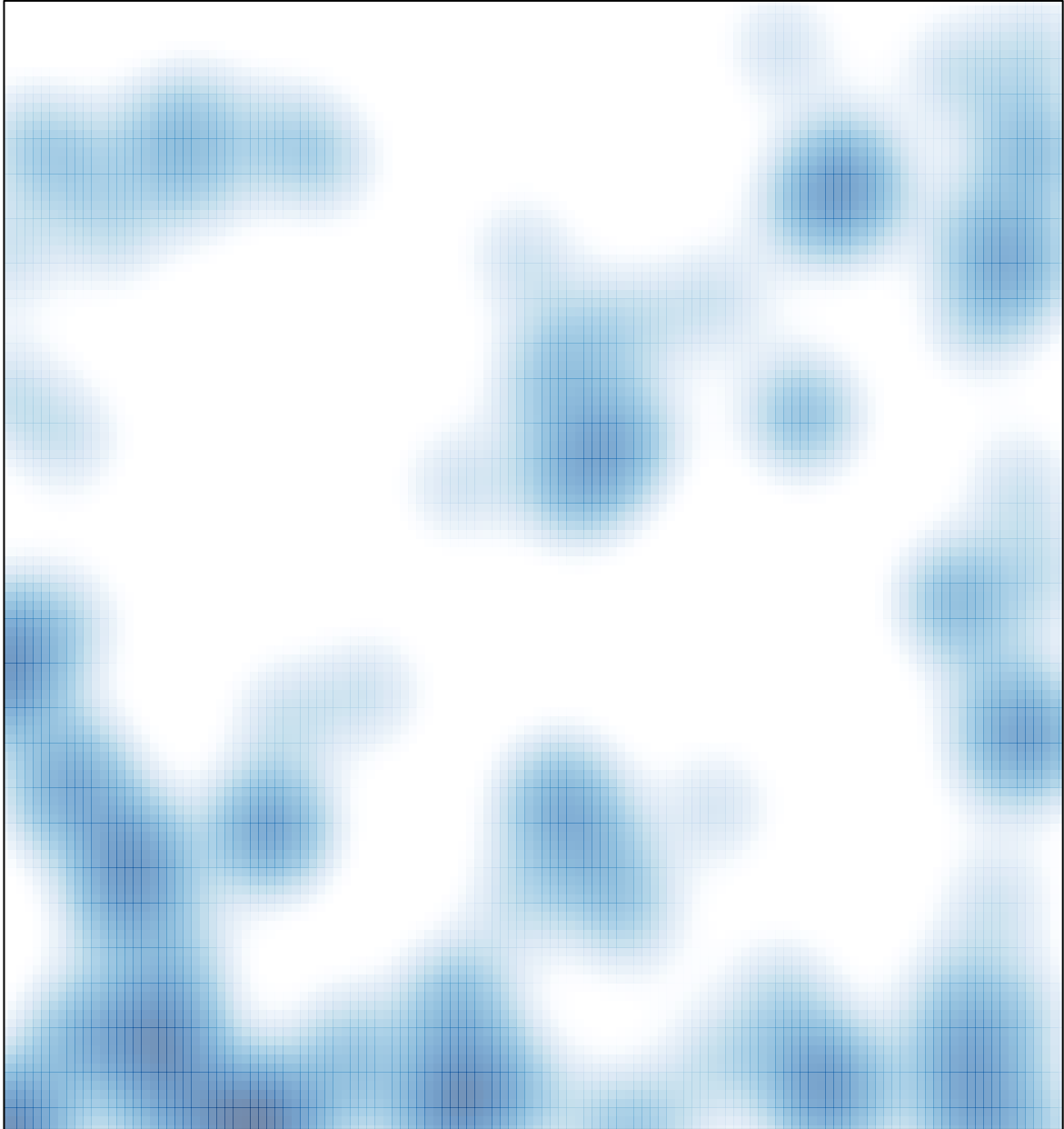


# protein dephosphorylation



# features = 130  
chi-square p = 0.75

# protein dephosphorylation



# features = 130 , max = 3