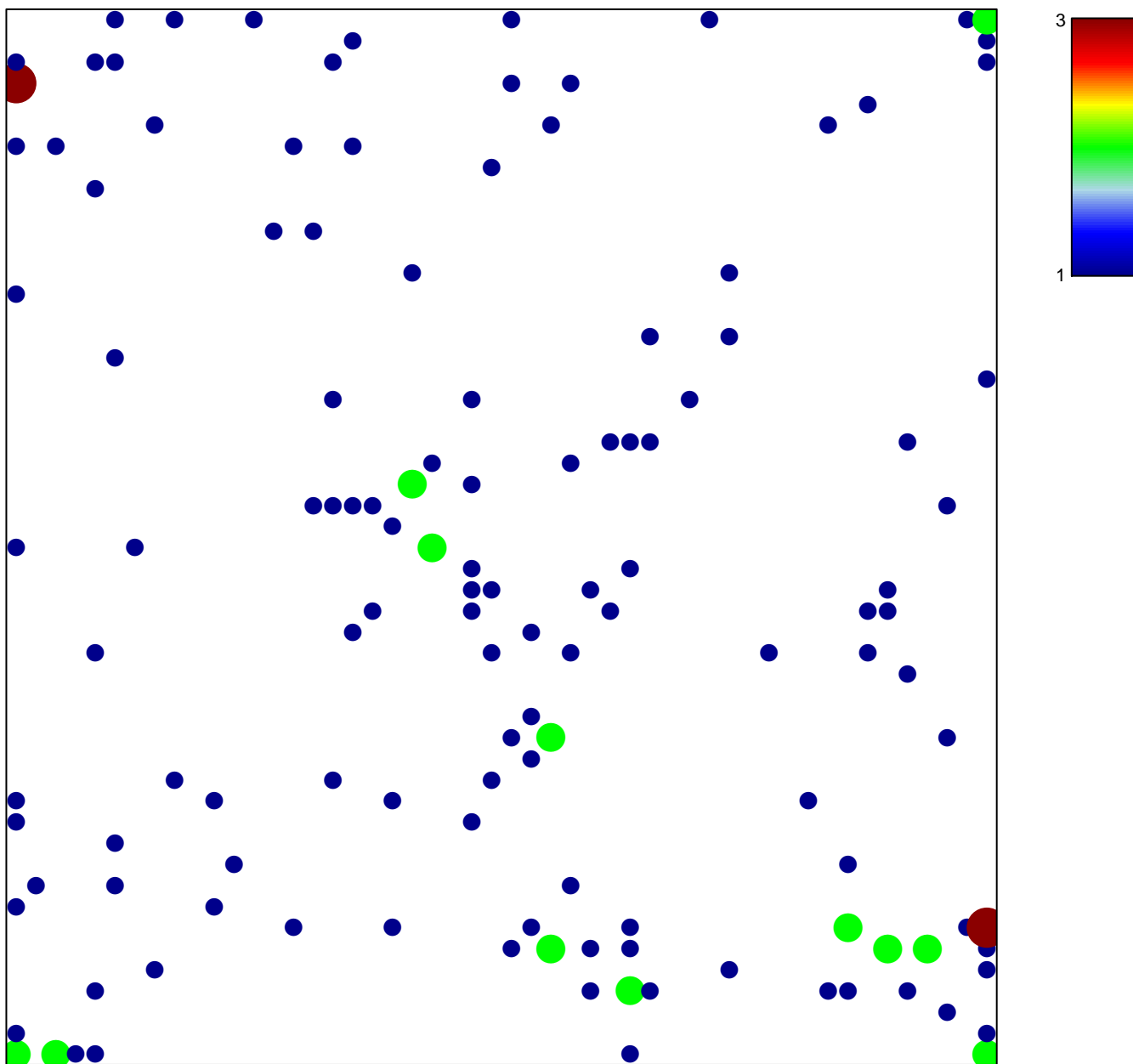
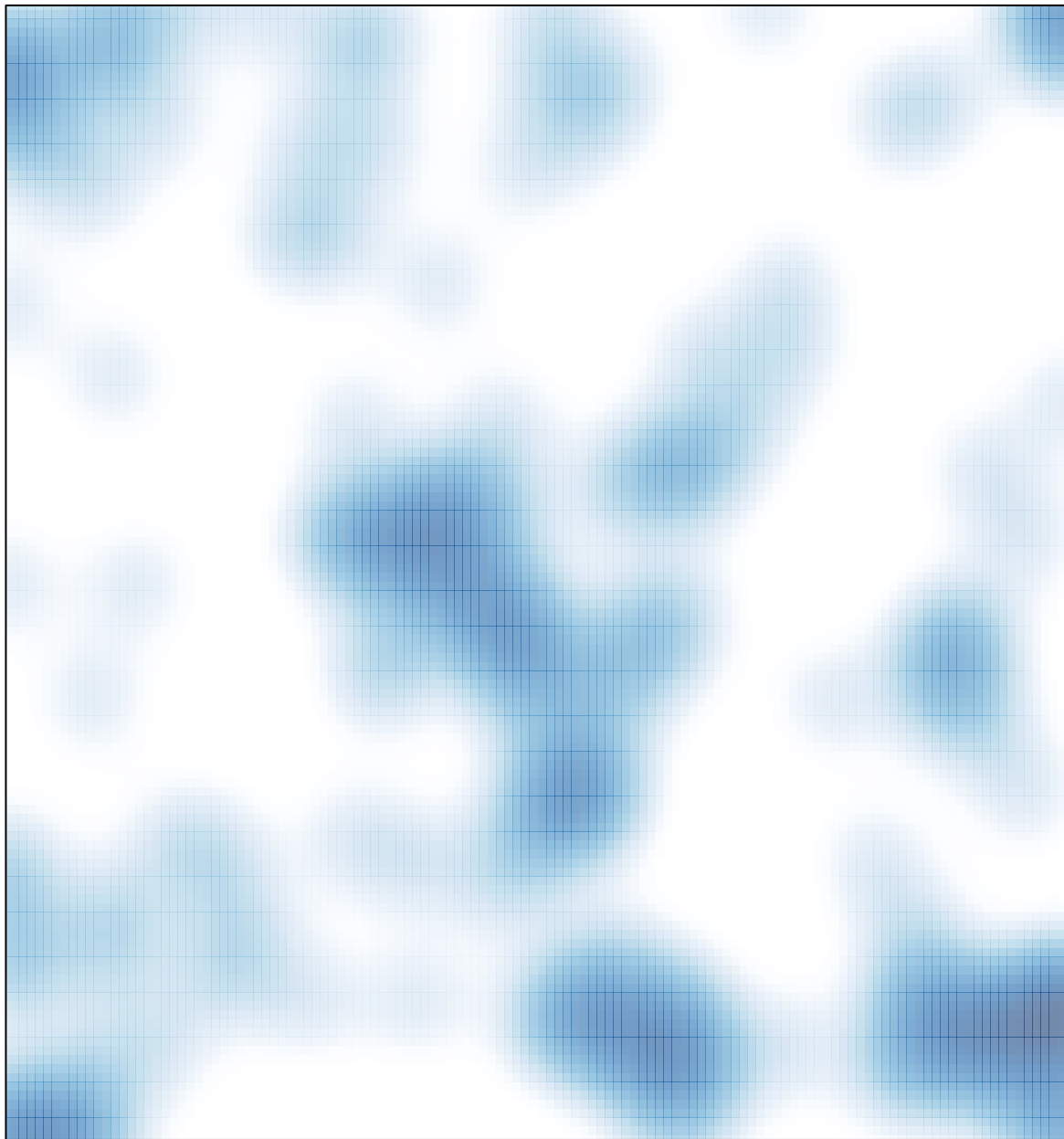


# carbohydrate binding



# features = 146  
chi-square p = 0.76

# carbohydrate binding



# features = 146 , max = 3