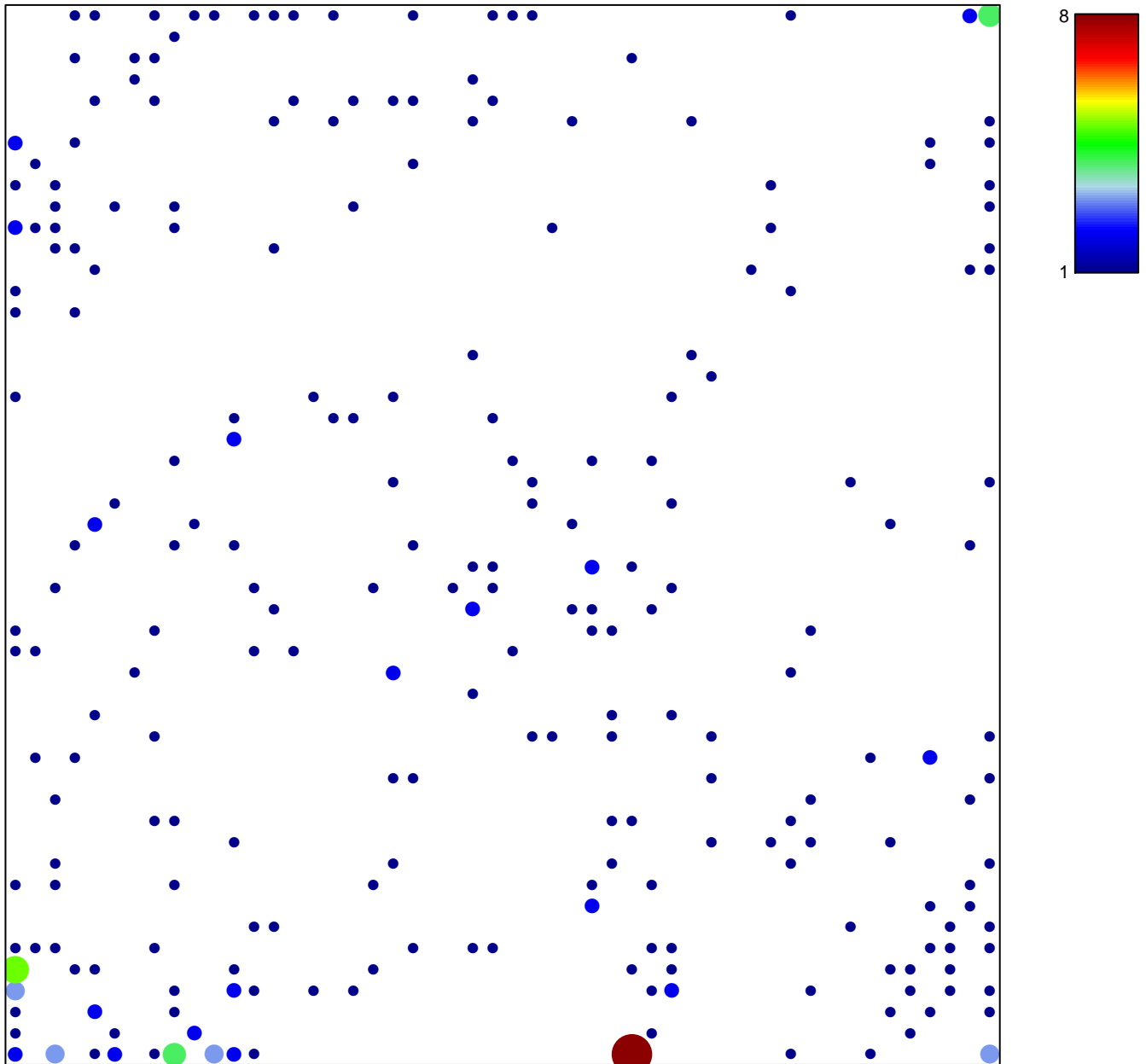
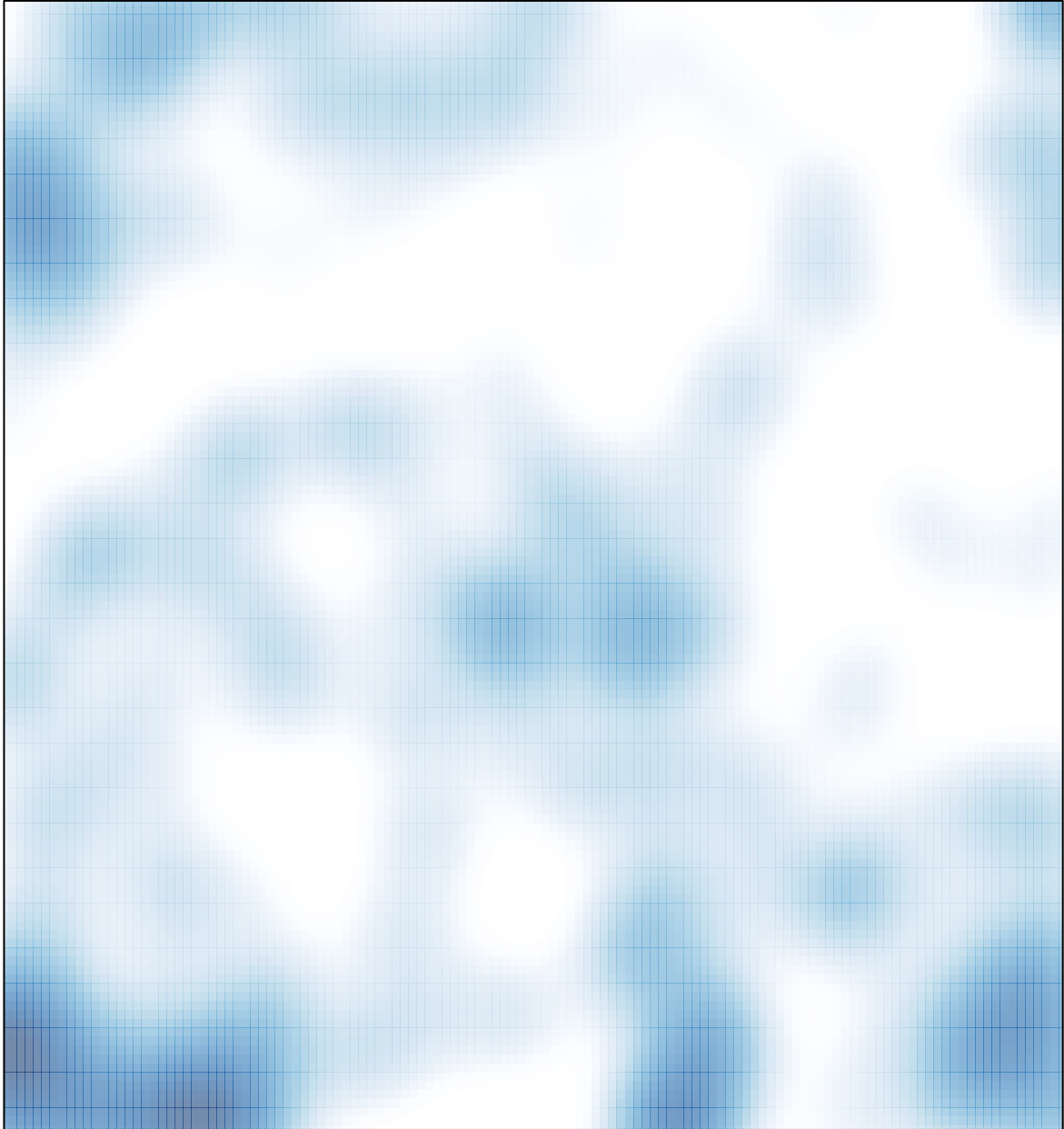


# receptor binding



# features = 281  
chi-square p = 0.68

# receptor binding



# features = 281 , max = 8