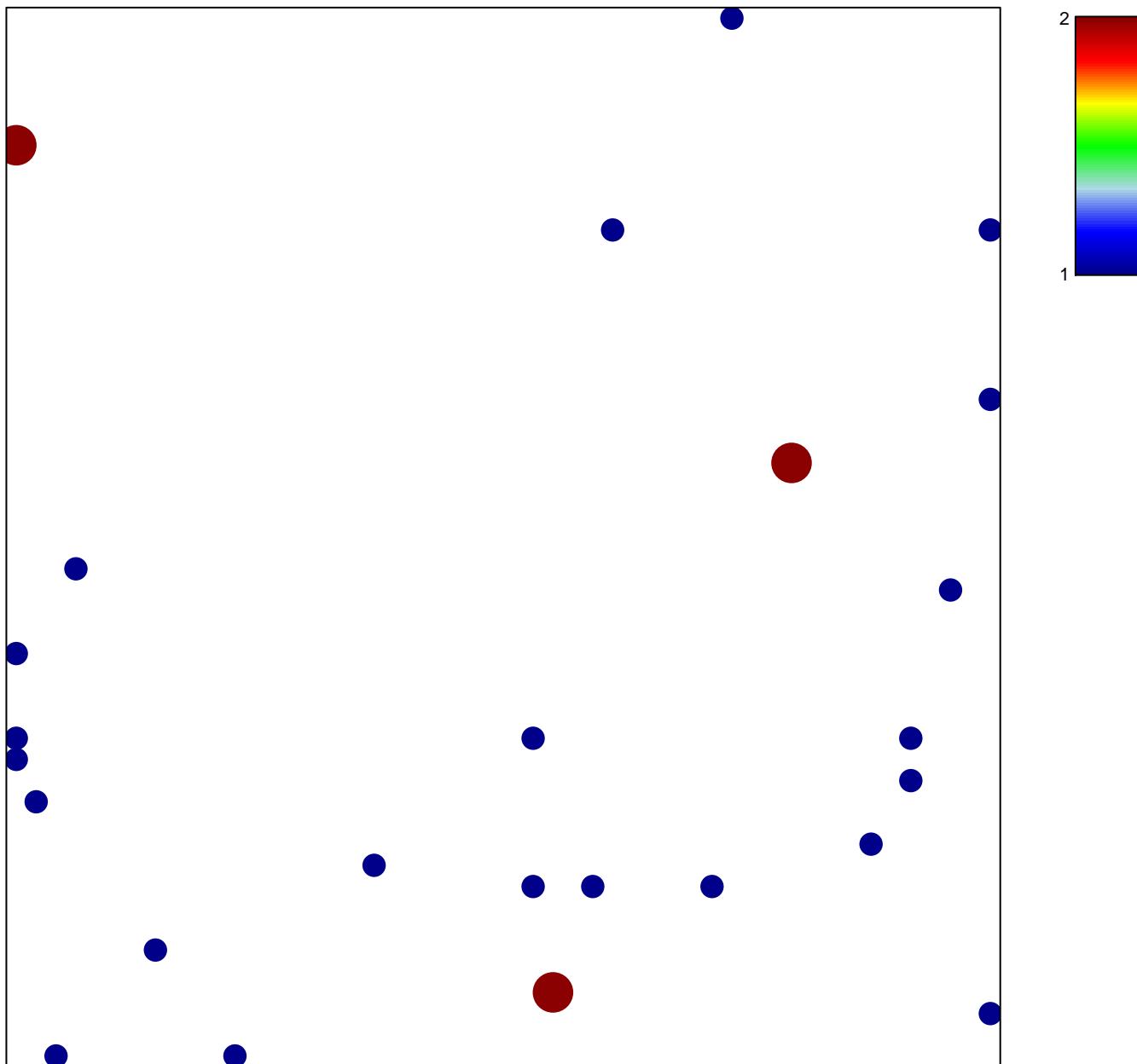
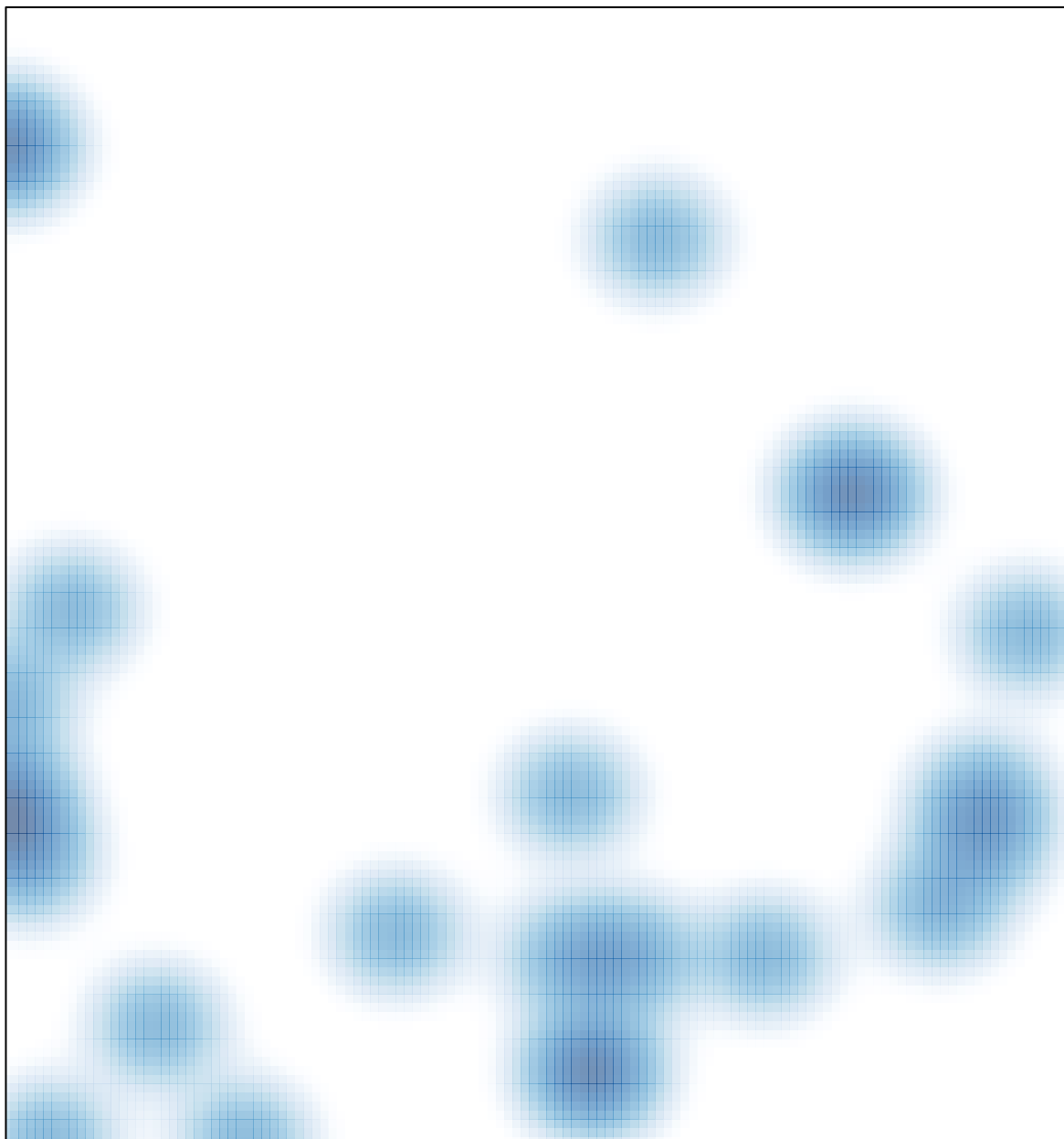


# cholesterol binding



# features = 28  
chi-square p = 0.81

# cholesterol binding



# features = 28 , max = 2