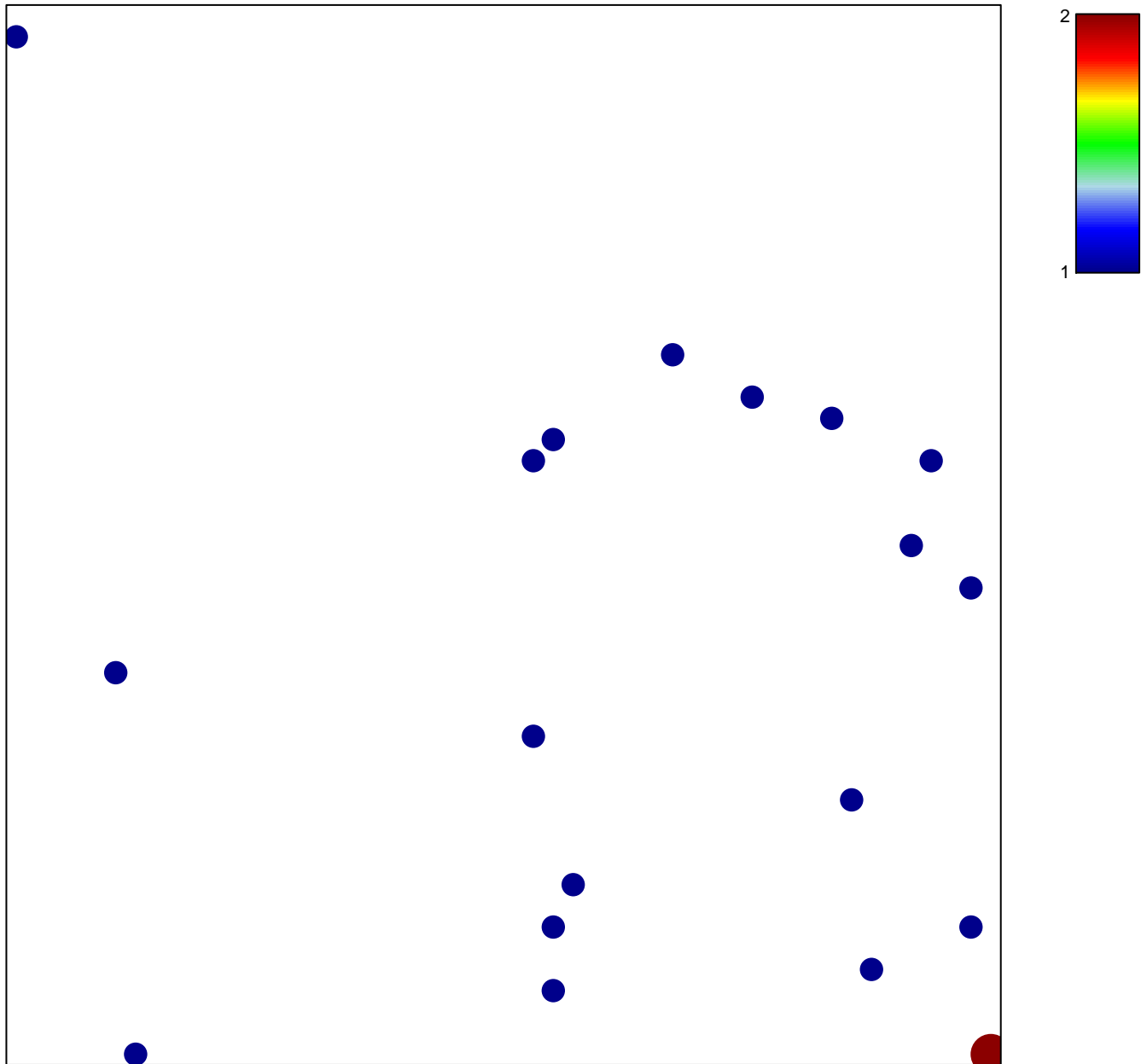
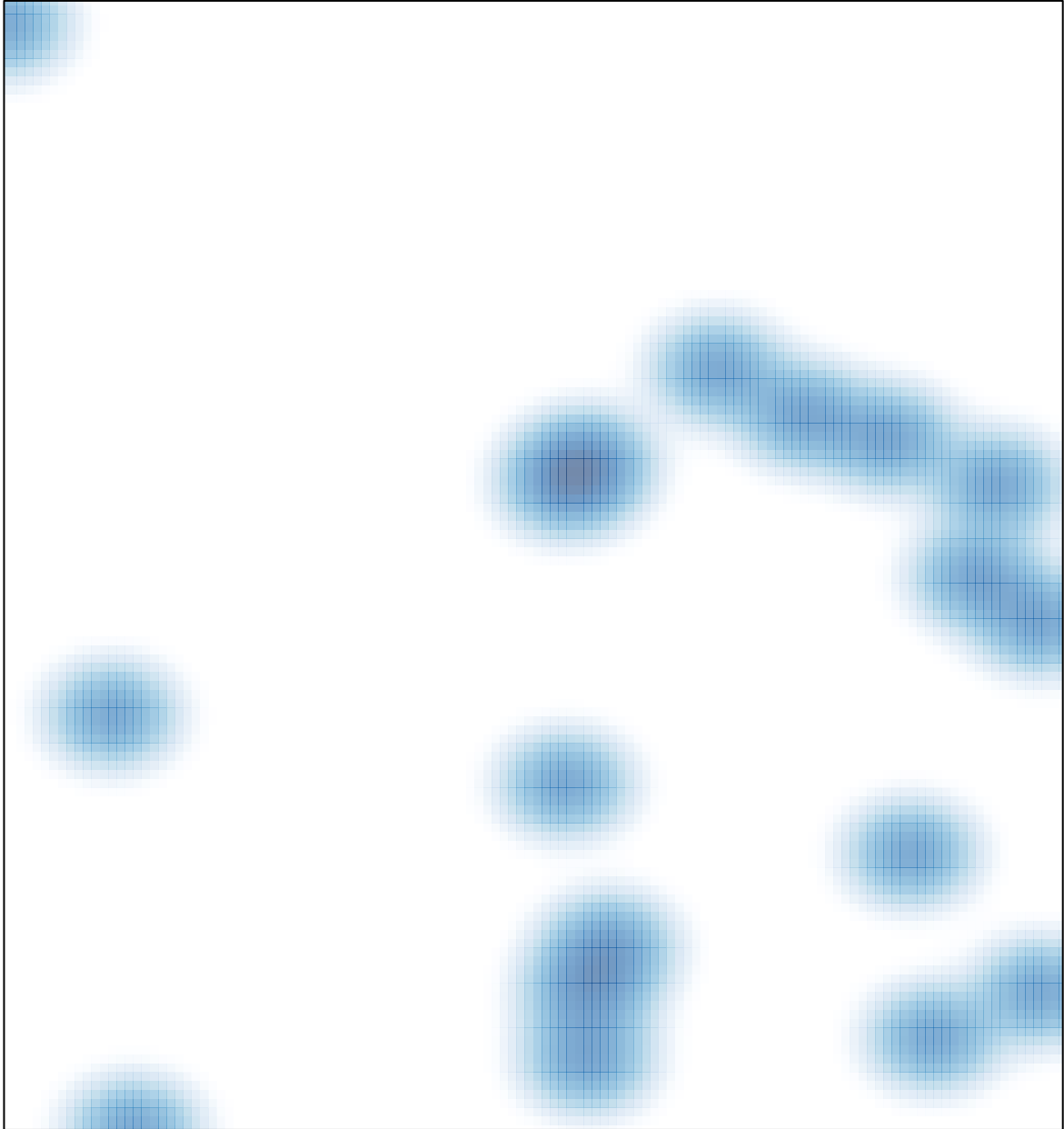


# NABA\_PROTEOGLYCANS



# features = 20  
chi-square p = 0.83

# NABA\_PROTEOGLYCANS



# features = 20 , max = 2