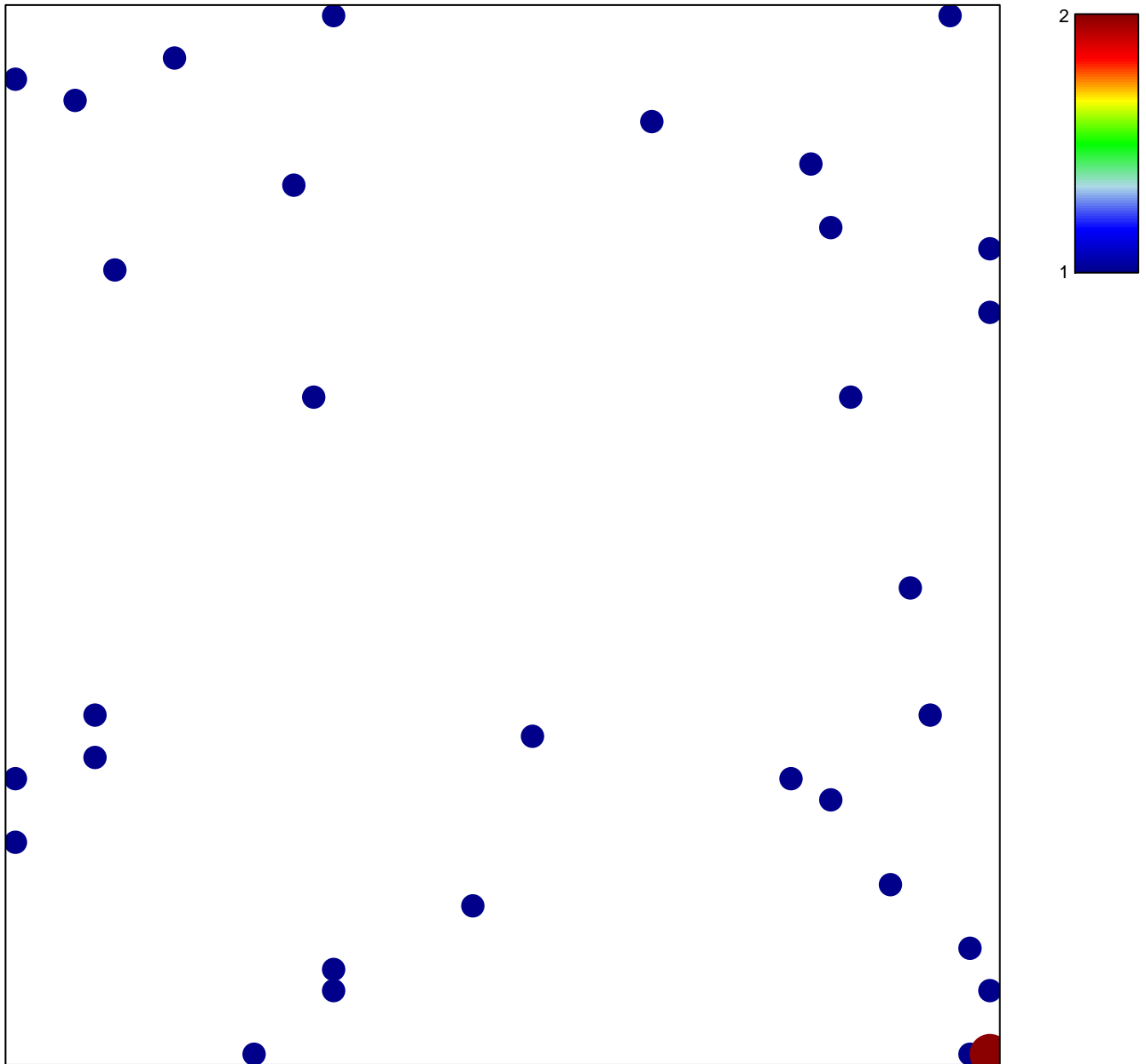
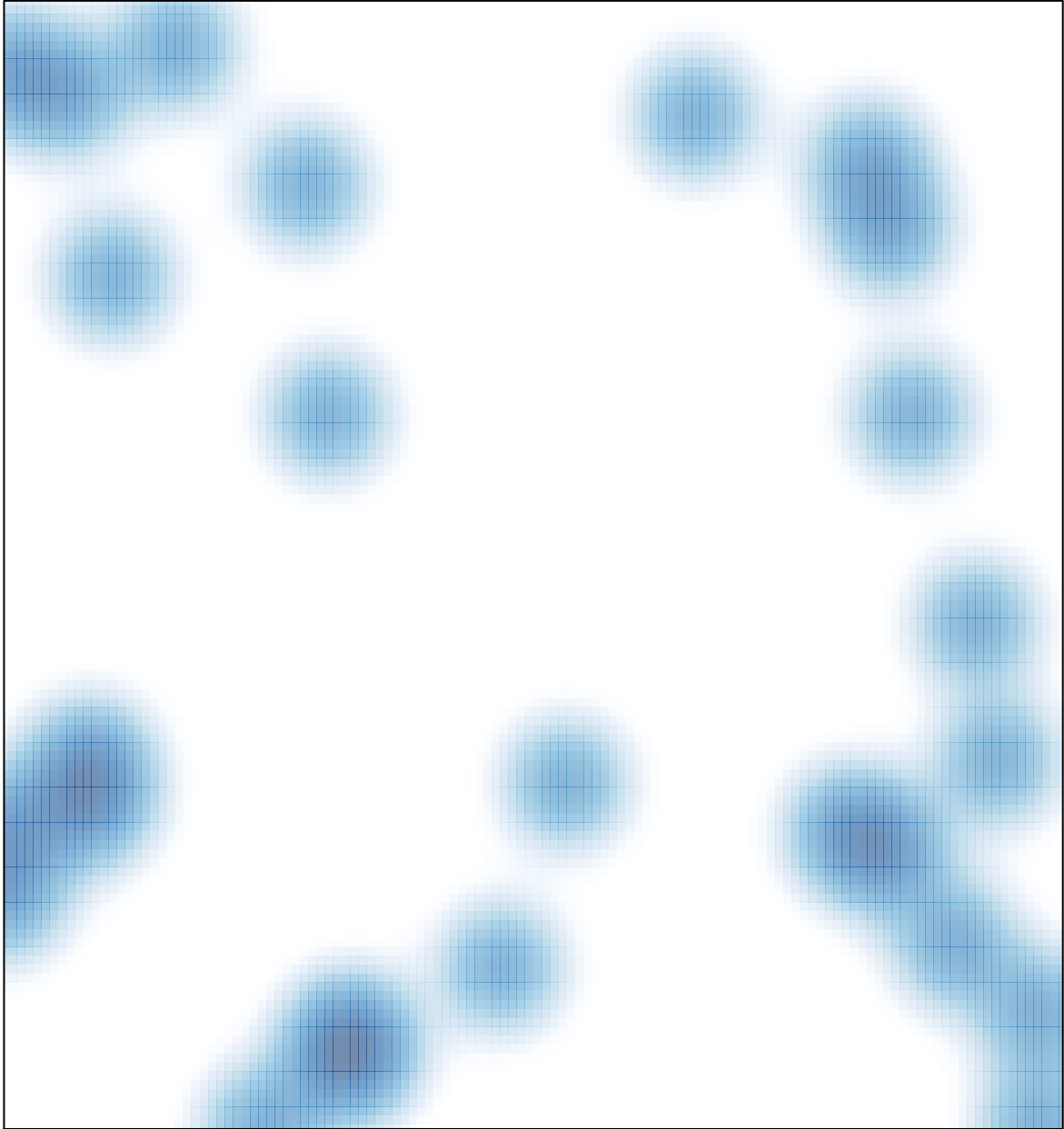


# SHAFFER\_IRF4\_MULTIPLE\_MYELOMA\_PROGRAM



# features = 33  
chi-square p = 0.82

# SHAFFER\_IRF4\_MULTIPLE\_MYELOMA\_PROGRAM



# features = 33 , max = 2