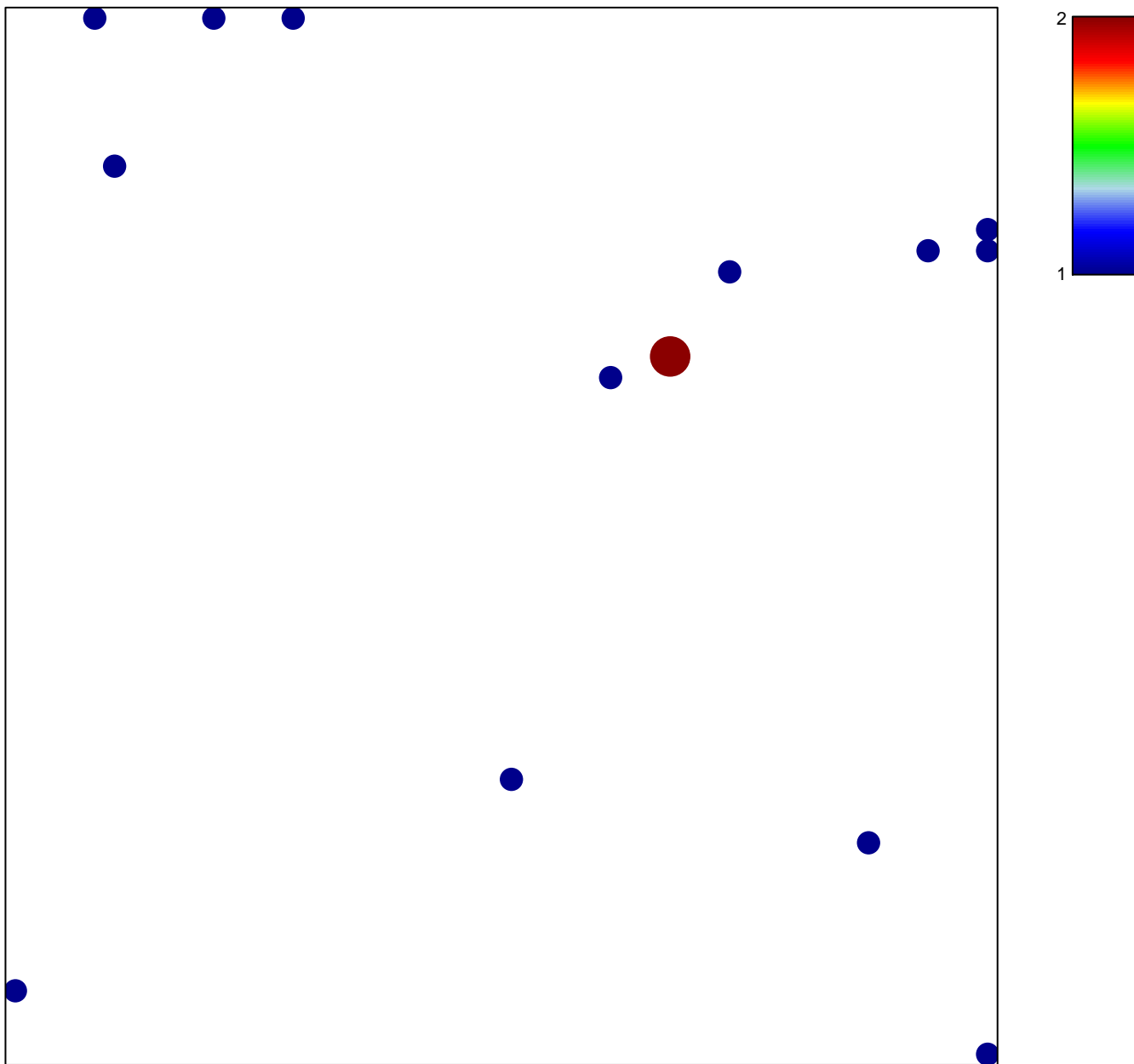
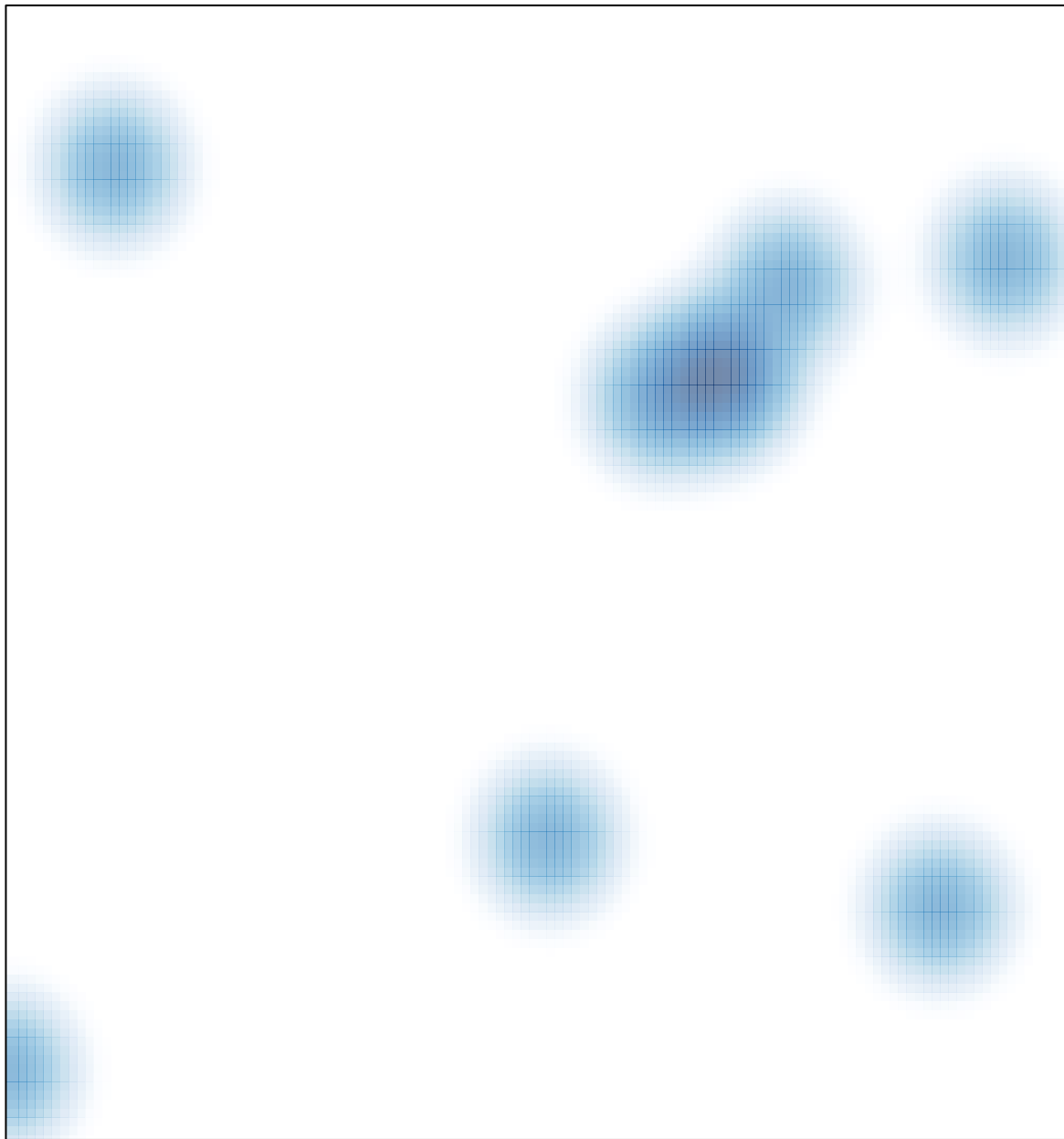


# MATTIOLI\_MULTIPLE\_MYELOMA\_SUBGROUPS



# features = 15  
chi-square p = 0.82

# MATTIOLI\_MULTIPLE\_MYELOMA\_SUBGROUPS



# features = 15 , max = 2