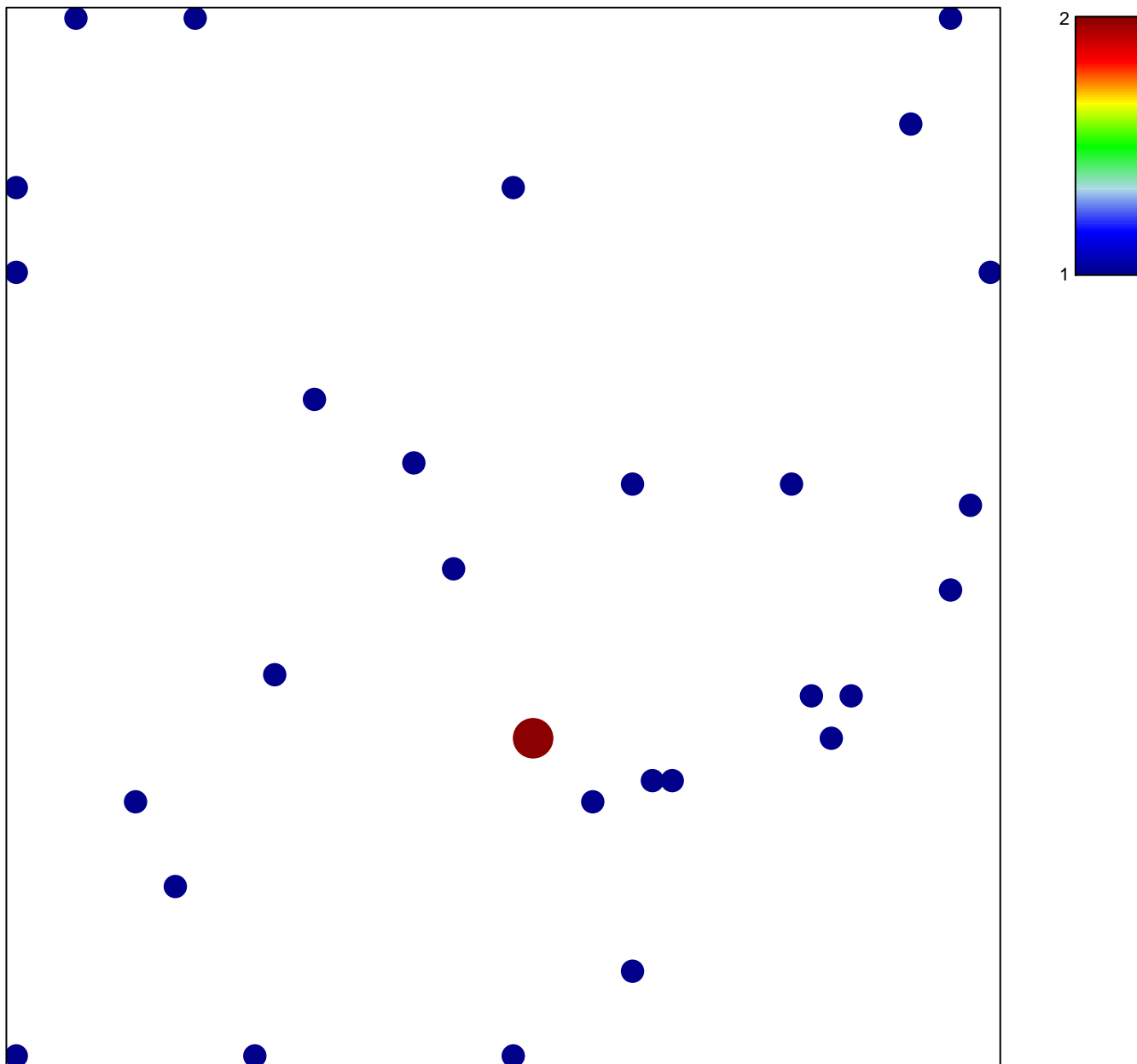
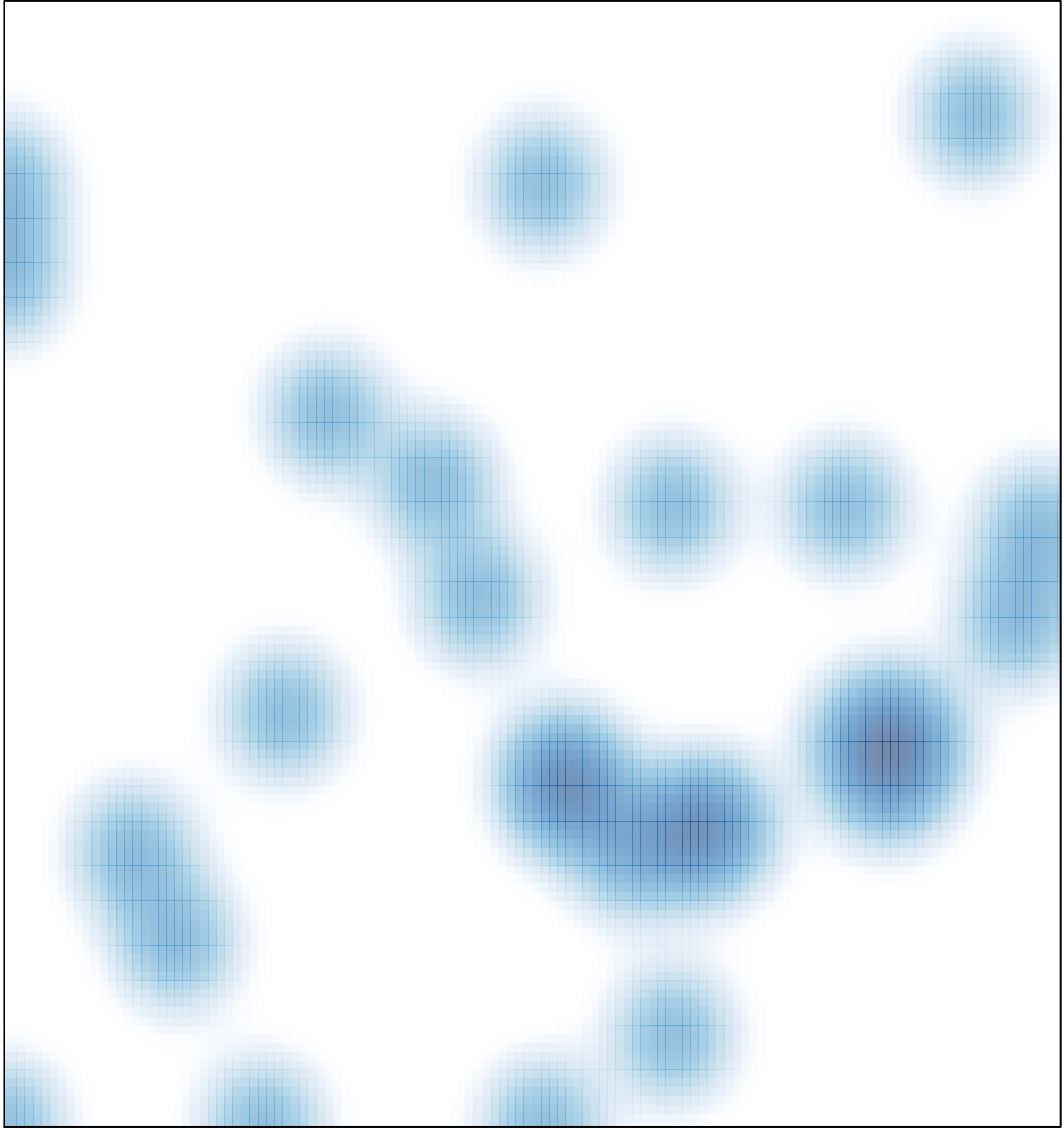


# MATTIOLI\_MULTIPLE\_MYELOMA\_WITH\_14Q32\_TRANSLOCATIONS



# features = 30  
chi-square p = 0.83

# DLI\_MULTIPLE\_MYELOMA\_WITH\_14Q32\_TRANSLOC



# features = 30 , max = 2