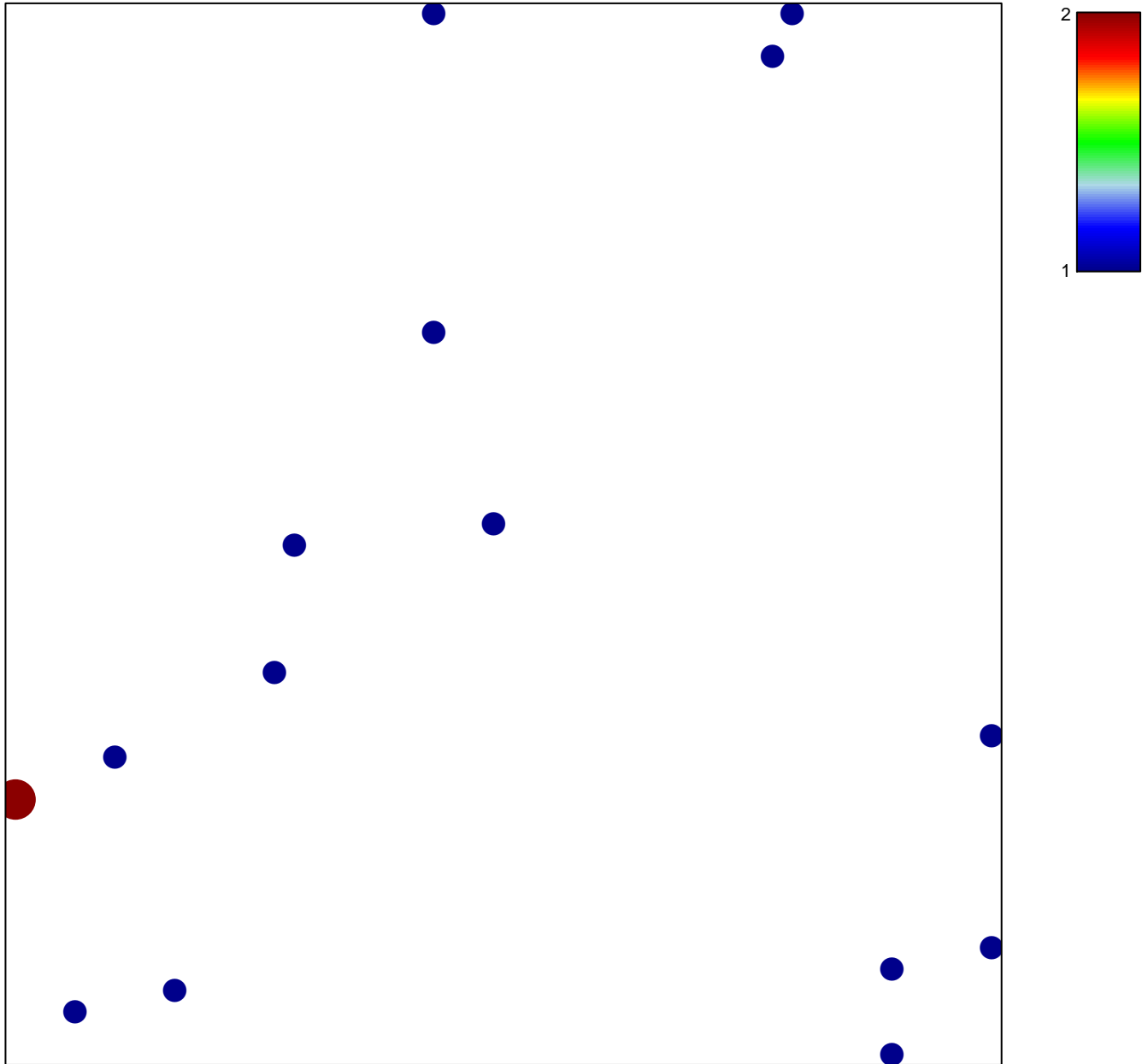
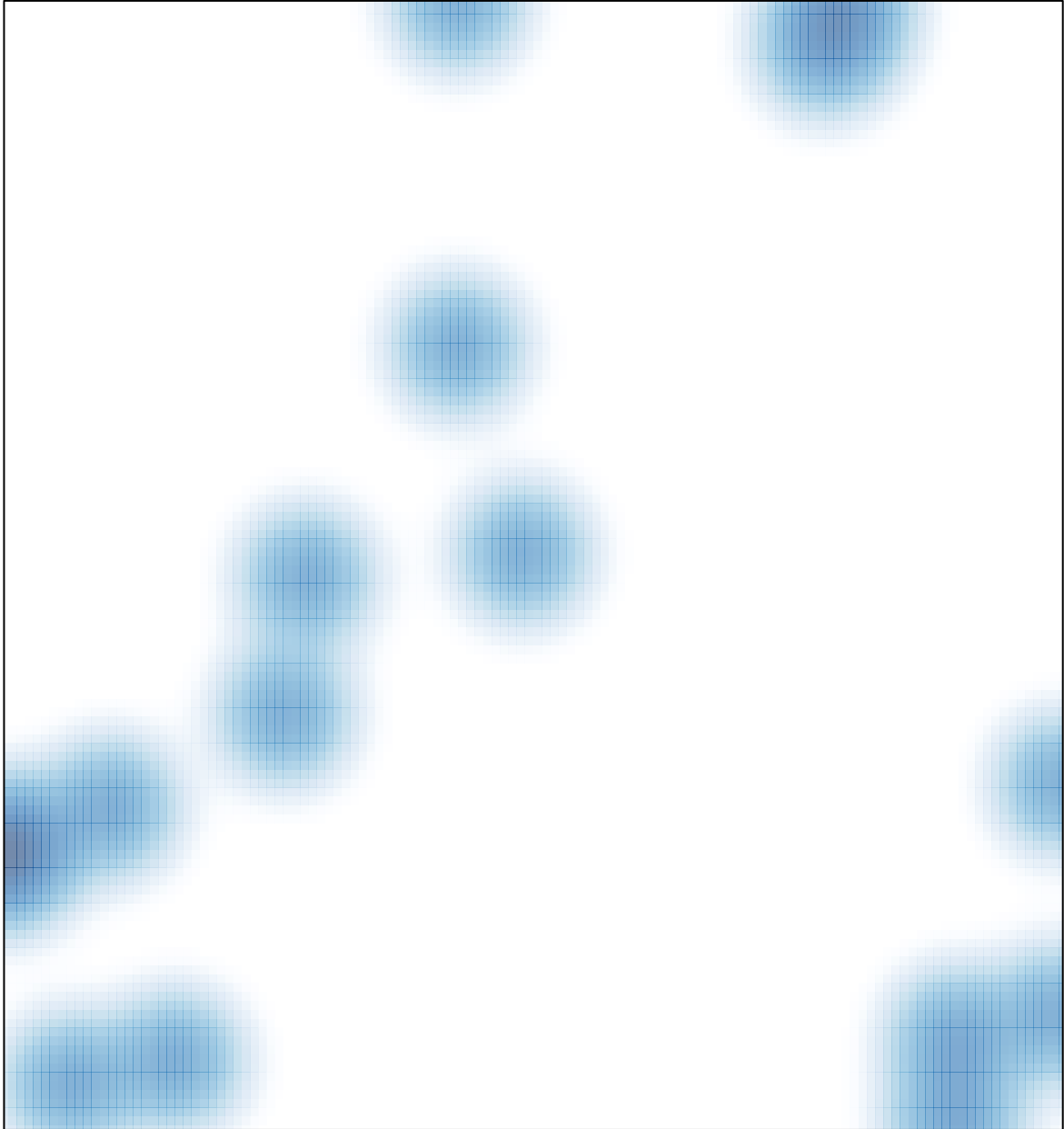


# ZHAN\_MULTIPLE\_MYELOMA\_HP\_DN



# features = 16  
chi-square p = 0.84

# ZHAN\_MULTIPLE\_MYELOMA\_HP\_DN



# features = 16 , max = 2