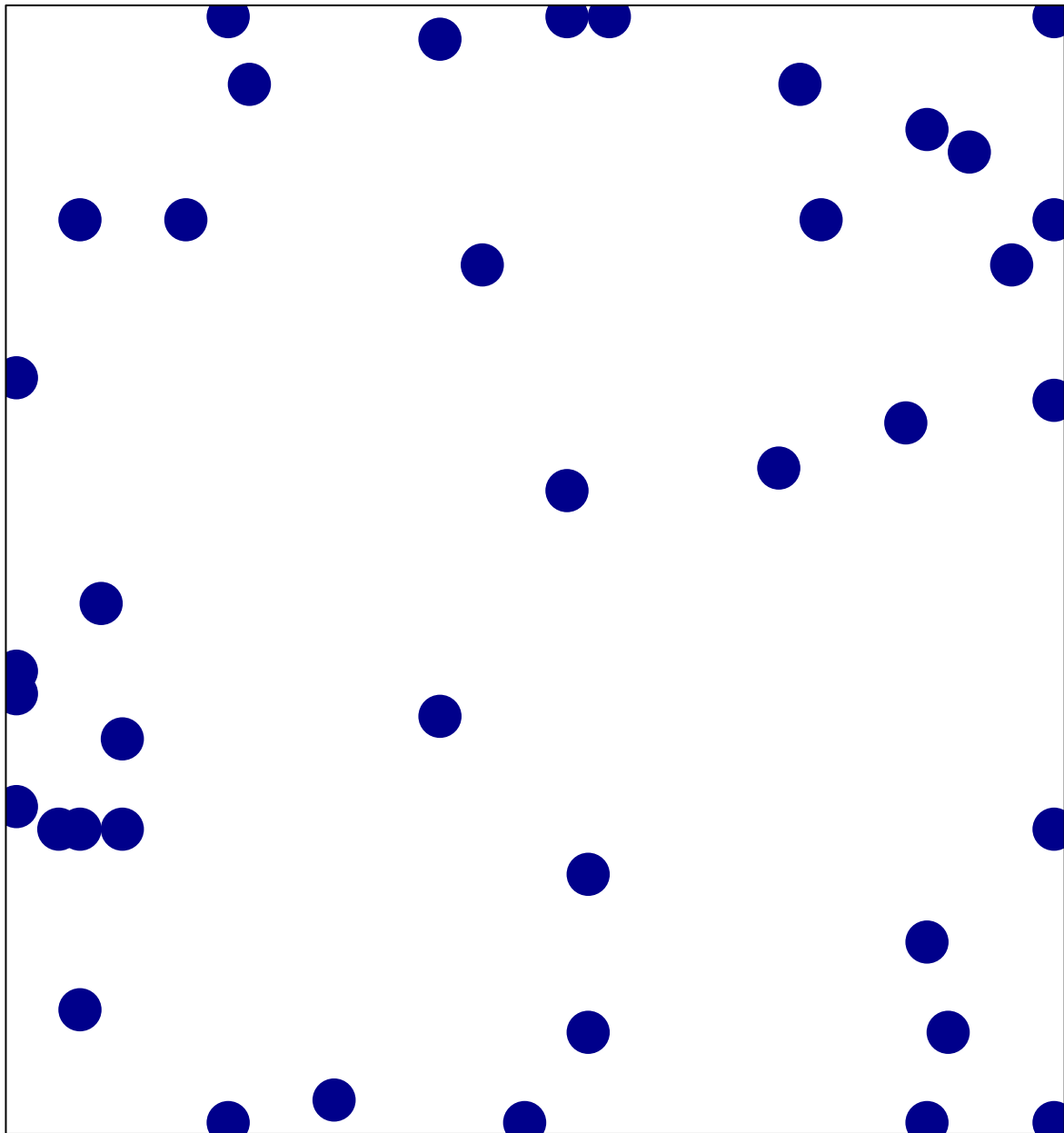
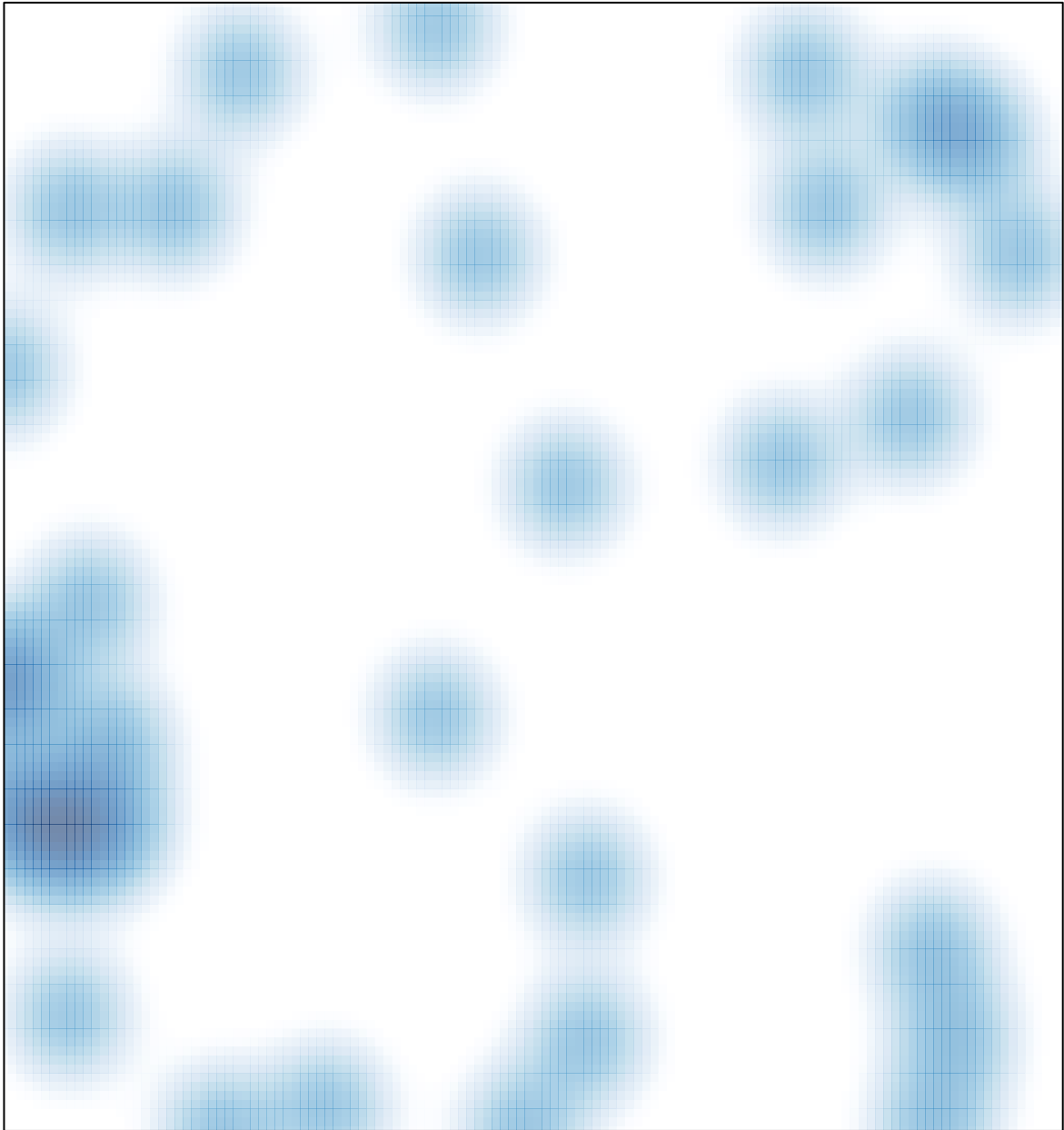


# ZHAN\_MULTIPLE\_MYELOMA\_MS\_DN



# features = 40  
chi-square p = 0.8

# ZHAN\_MULTIPLE\_MYELOMA\_MS\_DN



# features = 40 , max = 1