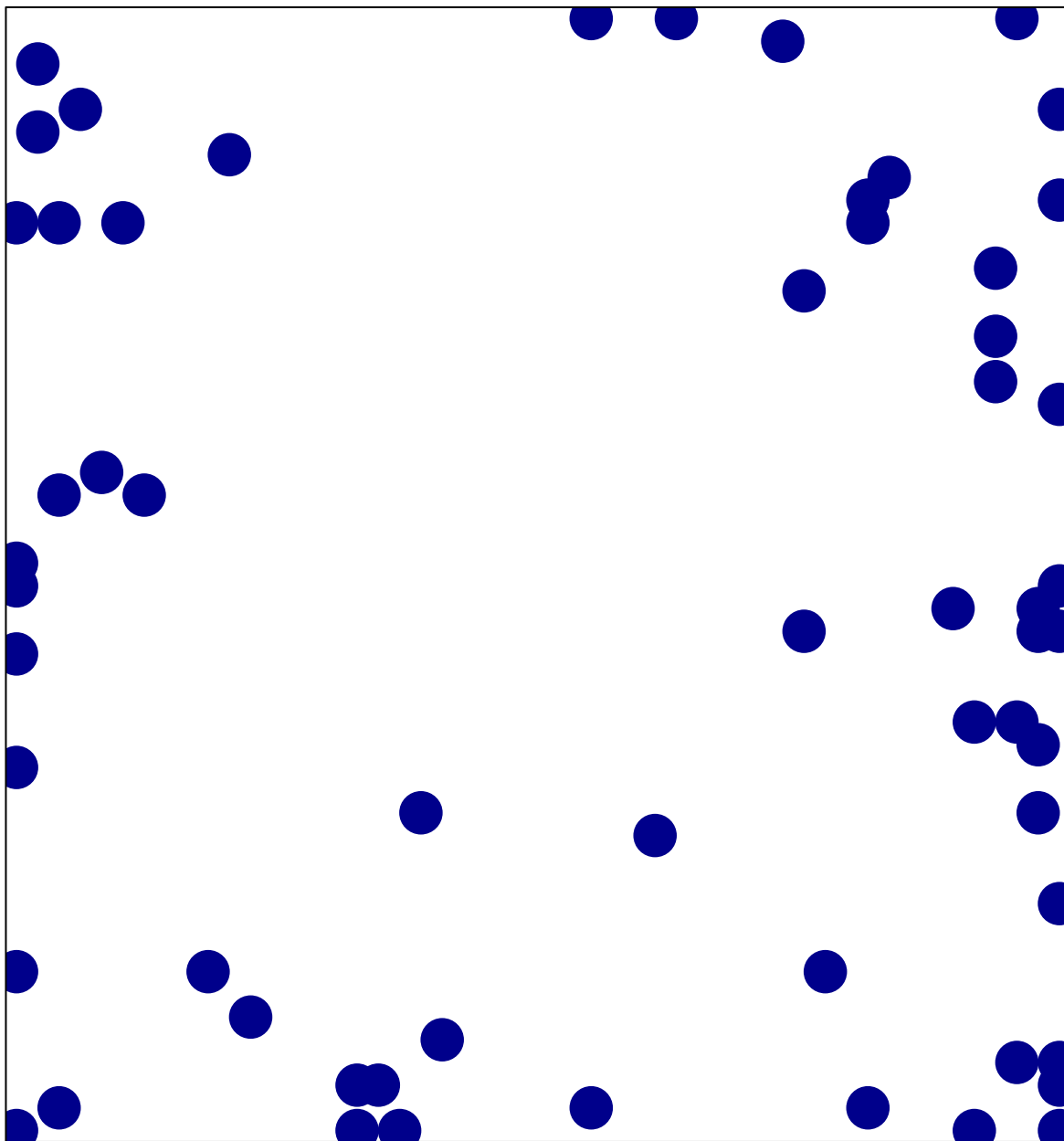
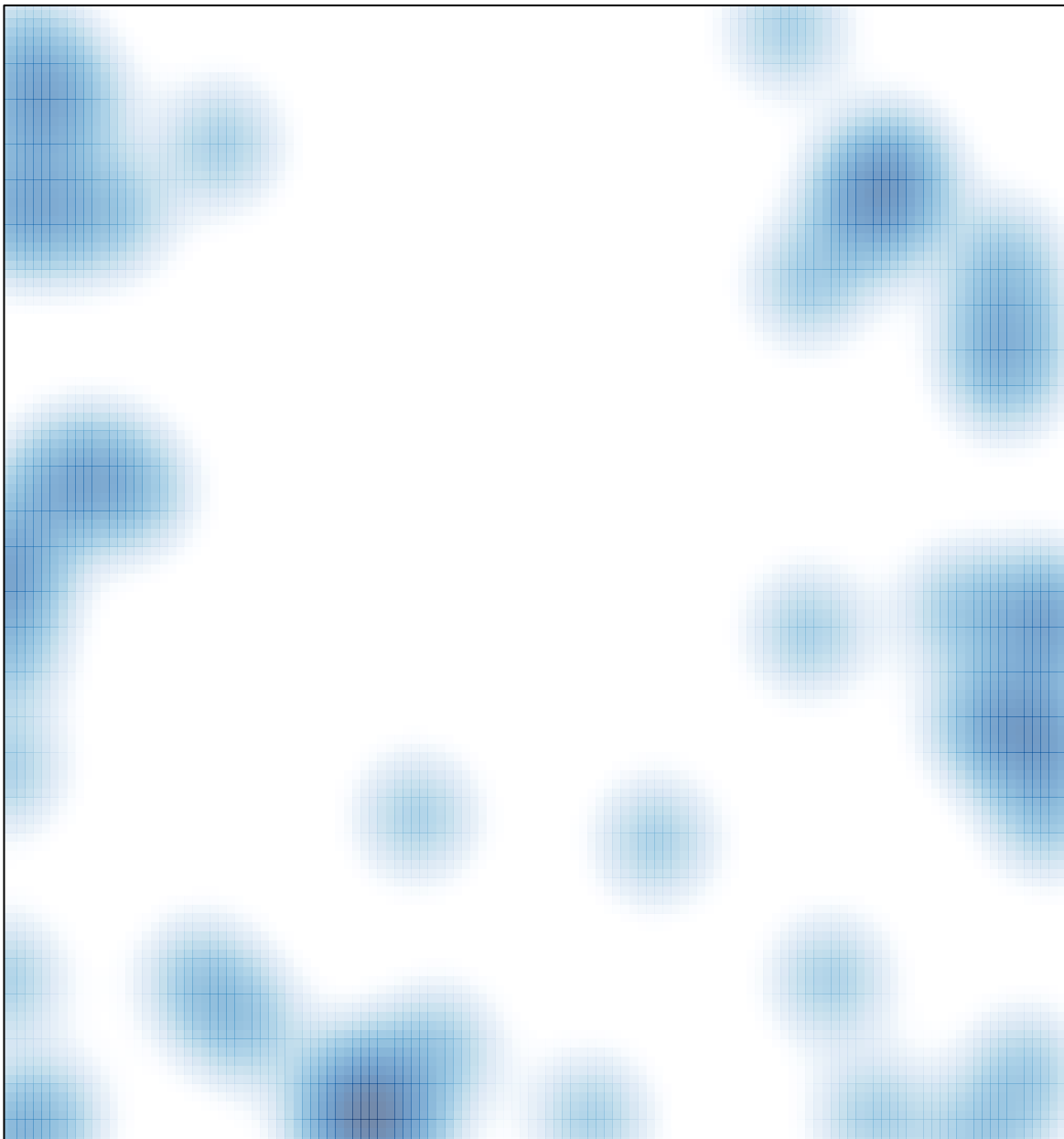


# ZHAN\_MULTIPLE\_MYELOMA\_UP



# features = 59  
chi-square p = 0.79

# ZHAN\_MULTIPLE\_MYELOMA\_UP



# features = 59 , max = 1