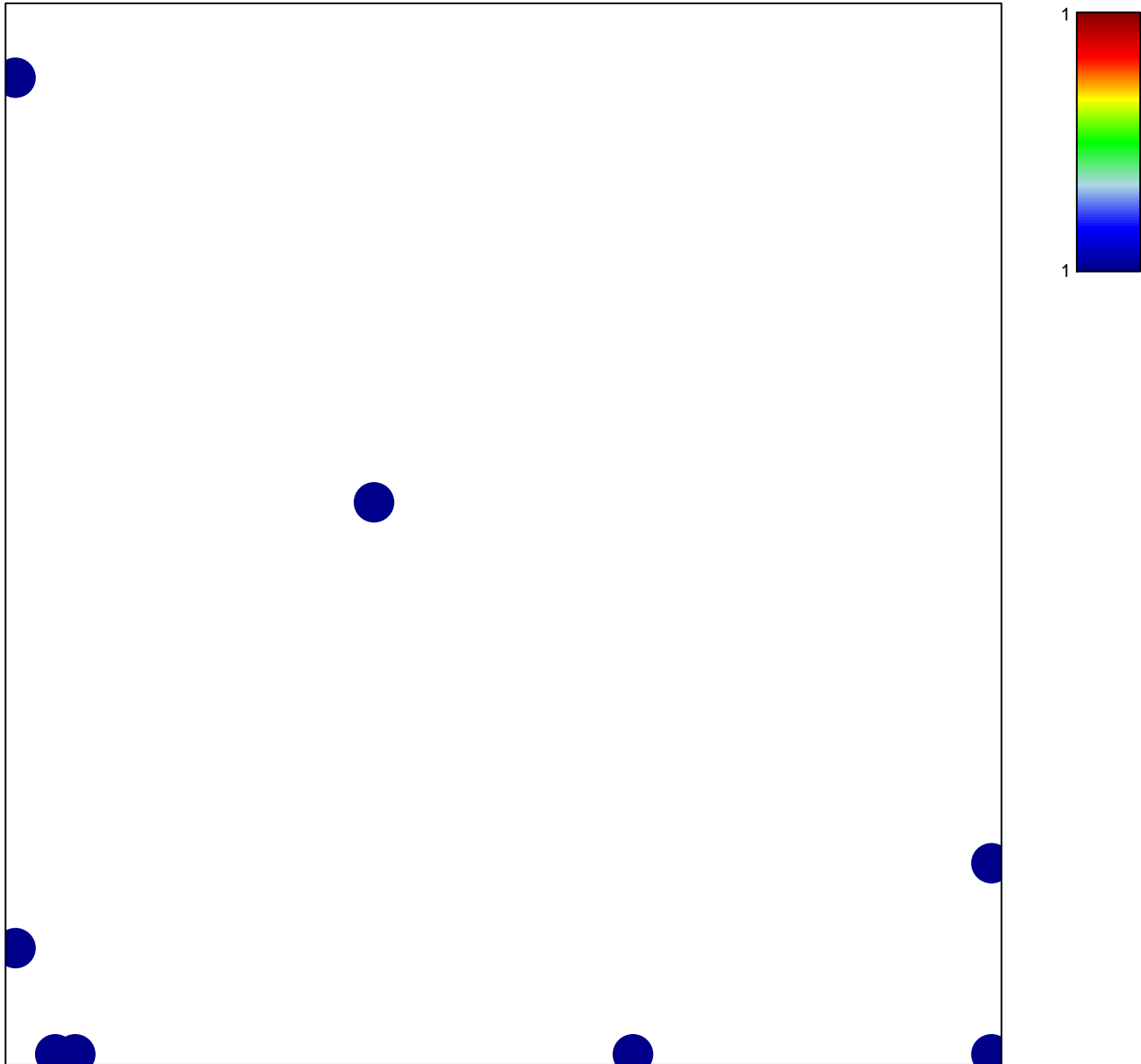
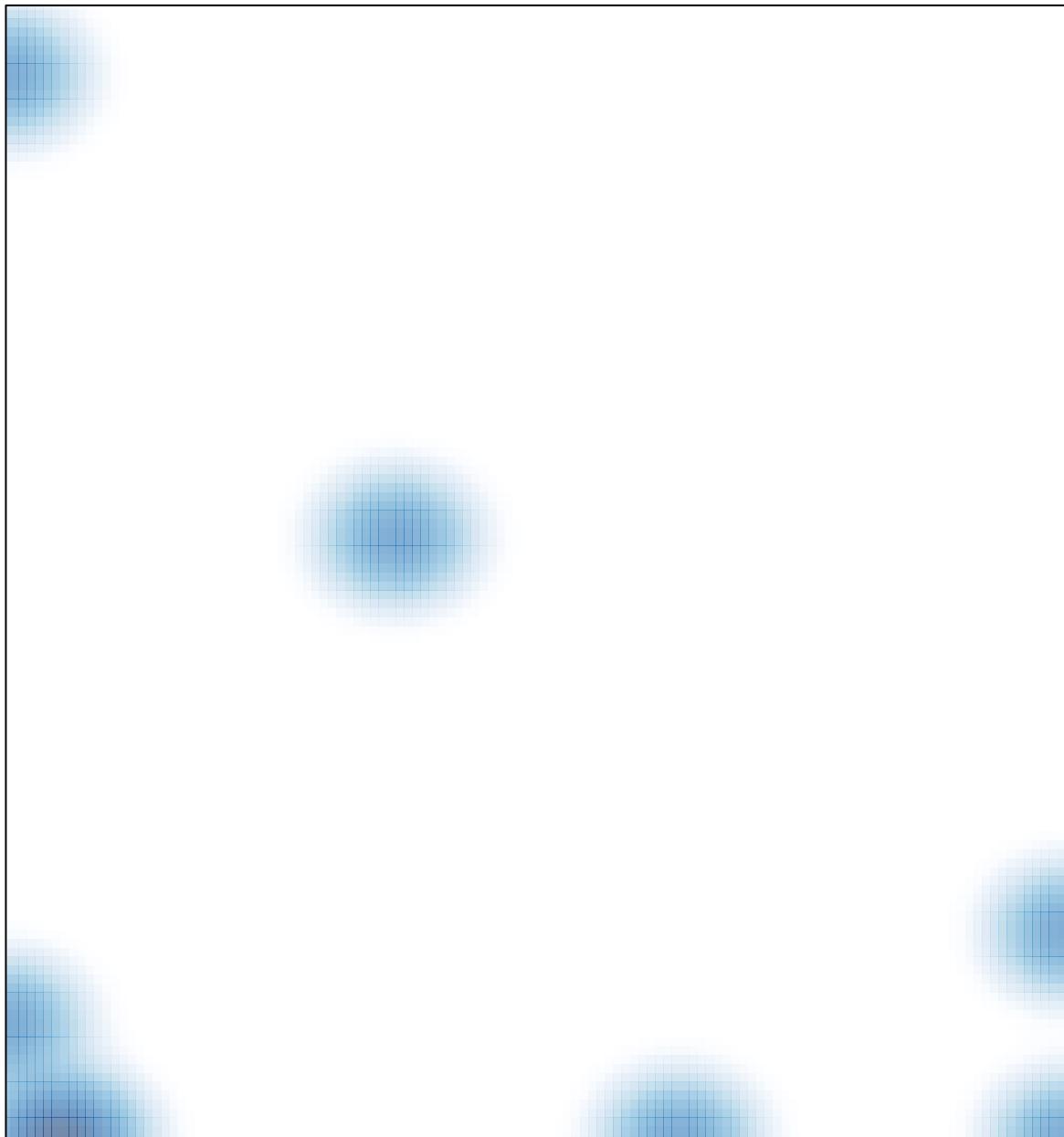


# ZHAN\_MULTIPLE\_MYELOMA\_SPIKED



# features = 8  
chi-square p = 0.79

# ZHAN\_MULTIPLE\_MYELOMA\_SPIKED



# features = 8 , max = 1