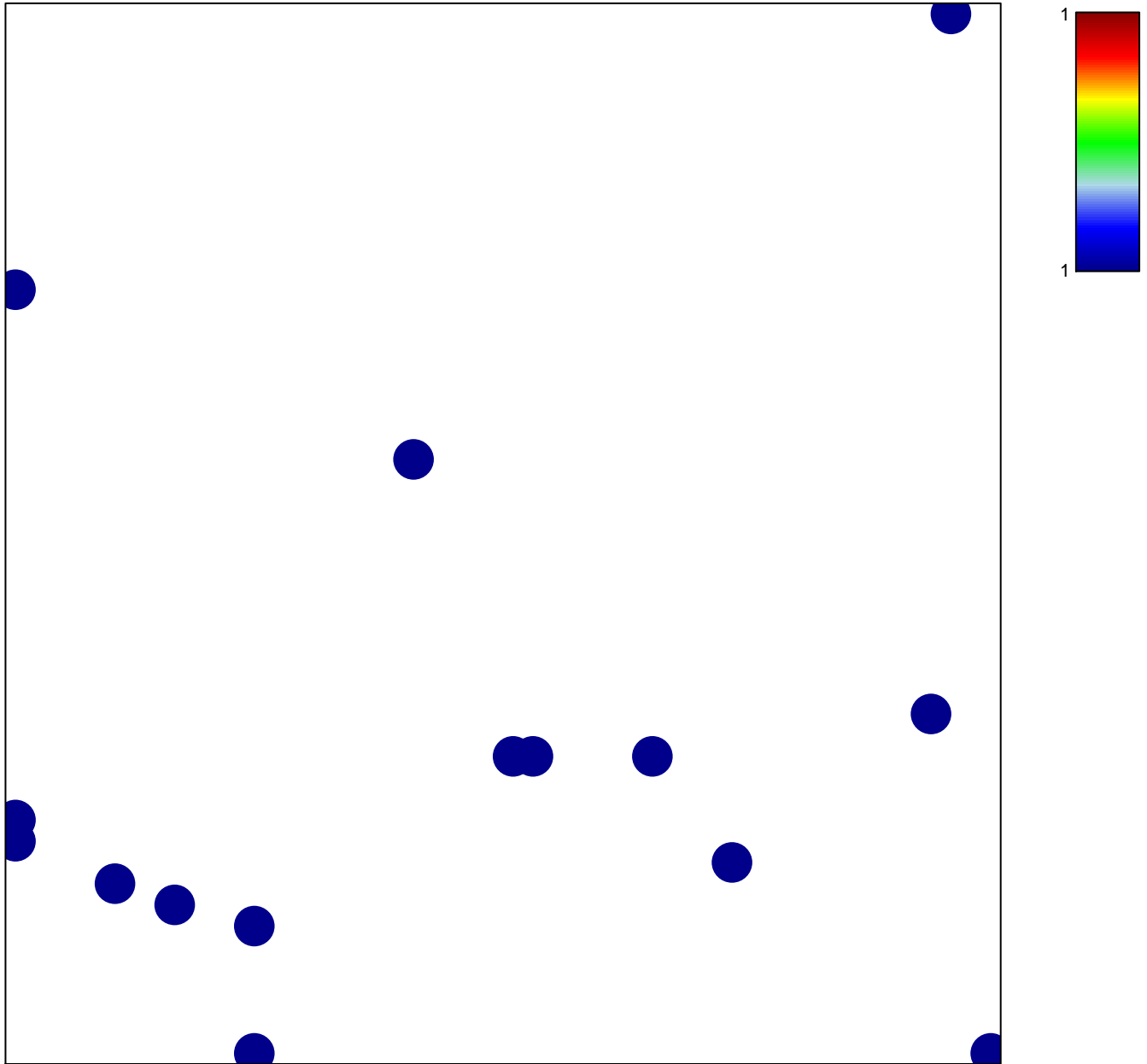


# ZHAN\_MULTIPLE\_MYELOMA\_SPIKED



# features = 15  
chi-square p = 0.82

# ZHAN\_MULTIPLE\_MYELOMA\_SPIKED



# features = 15 , max = 1