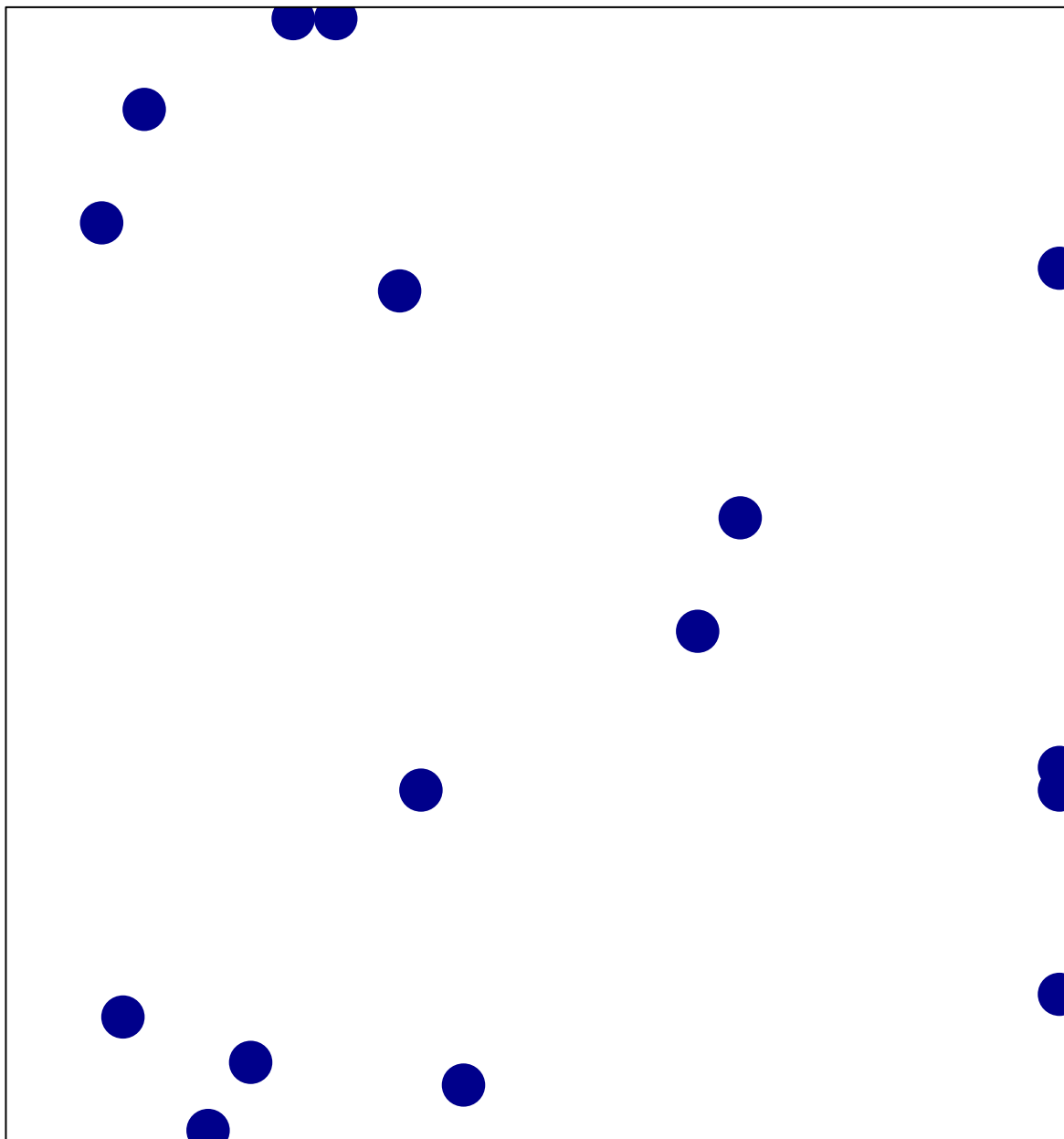
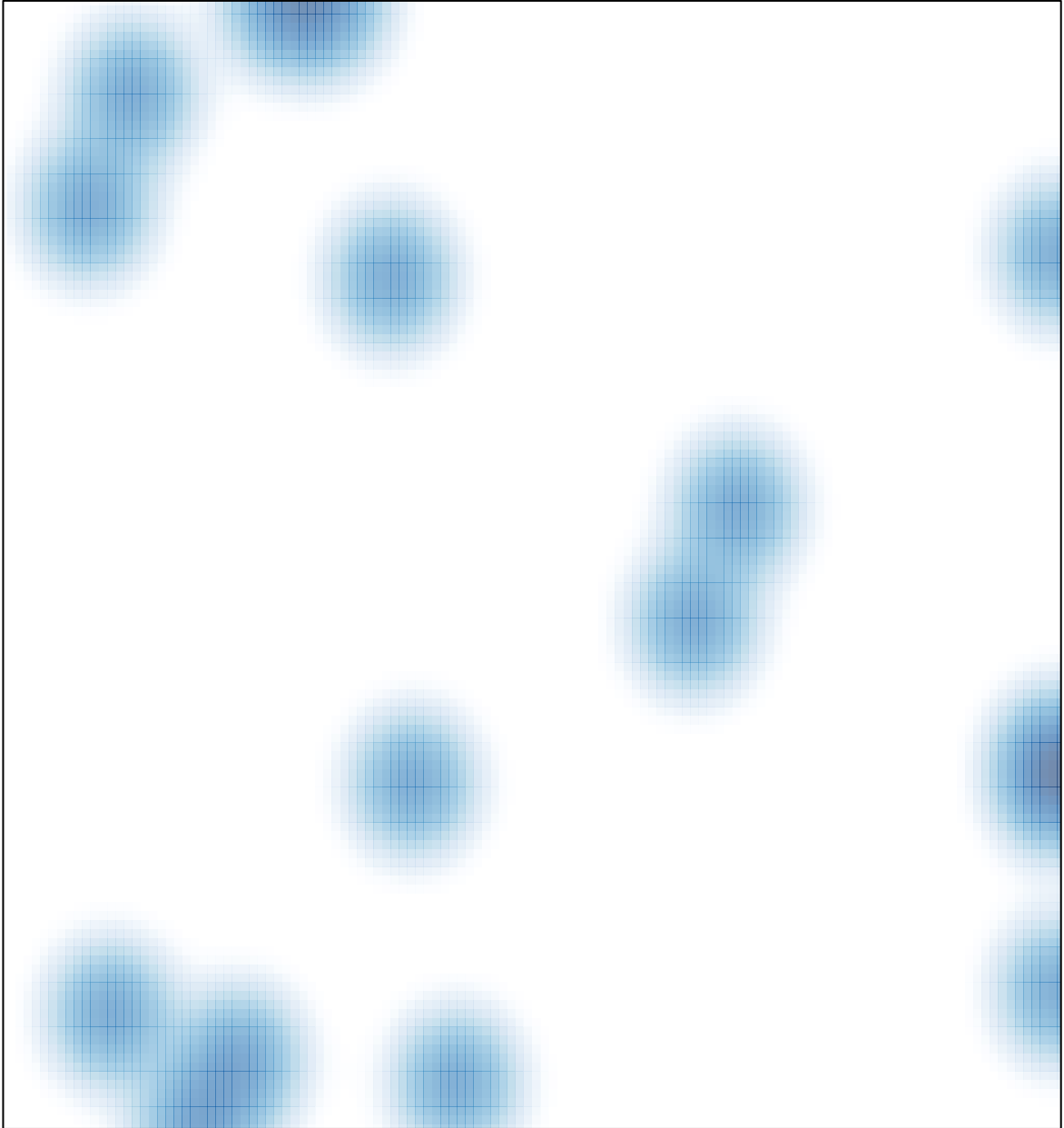


# ZHAN\_MULTIPLE\_MYELOMA\_MF\_DN



# features = 16  
chi-square p = 0.83

# ZHAN\_MULTIPLE\_MYELOMA\_MF\_DN



# features = 16 , max = 1