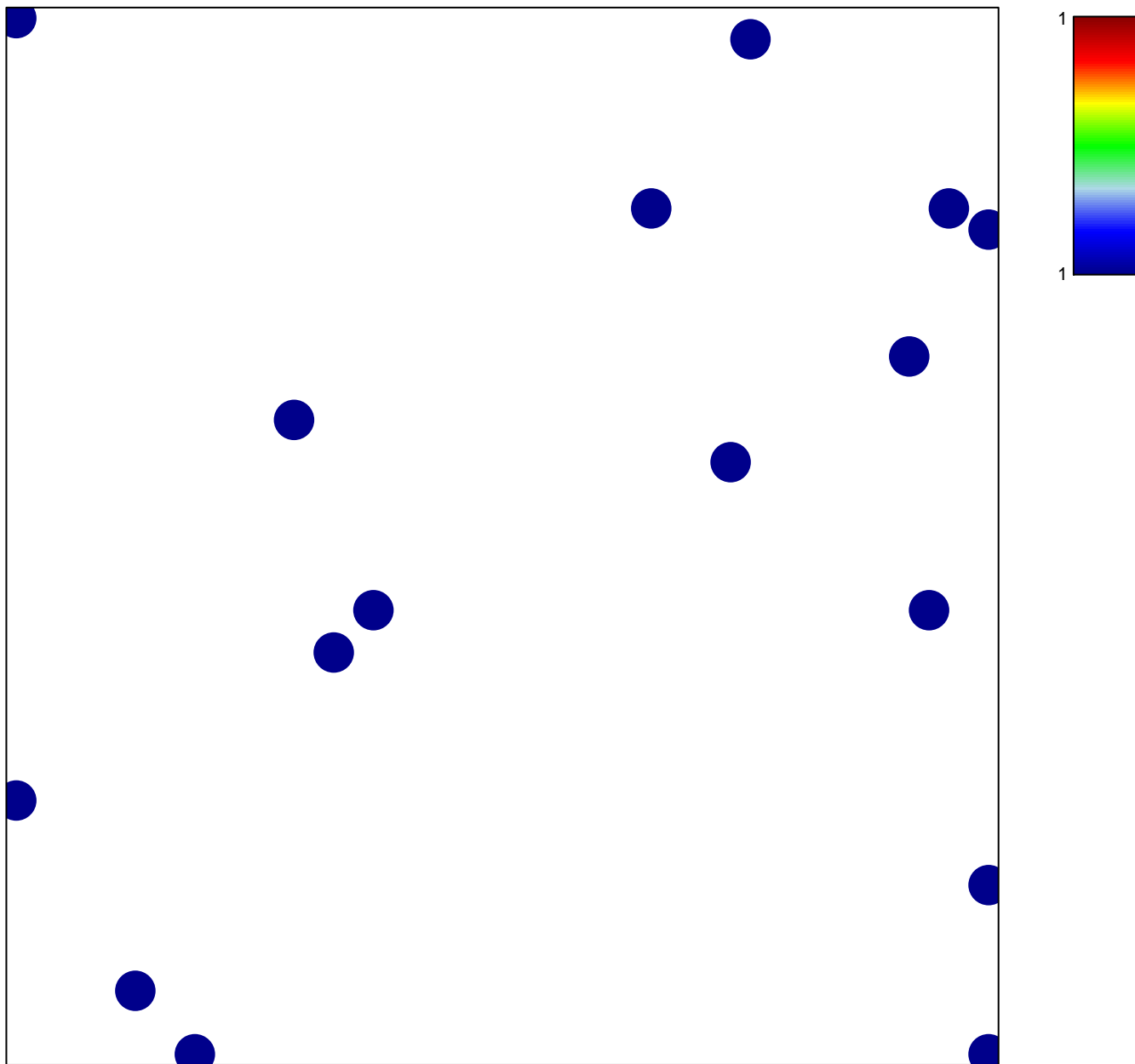
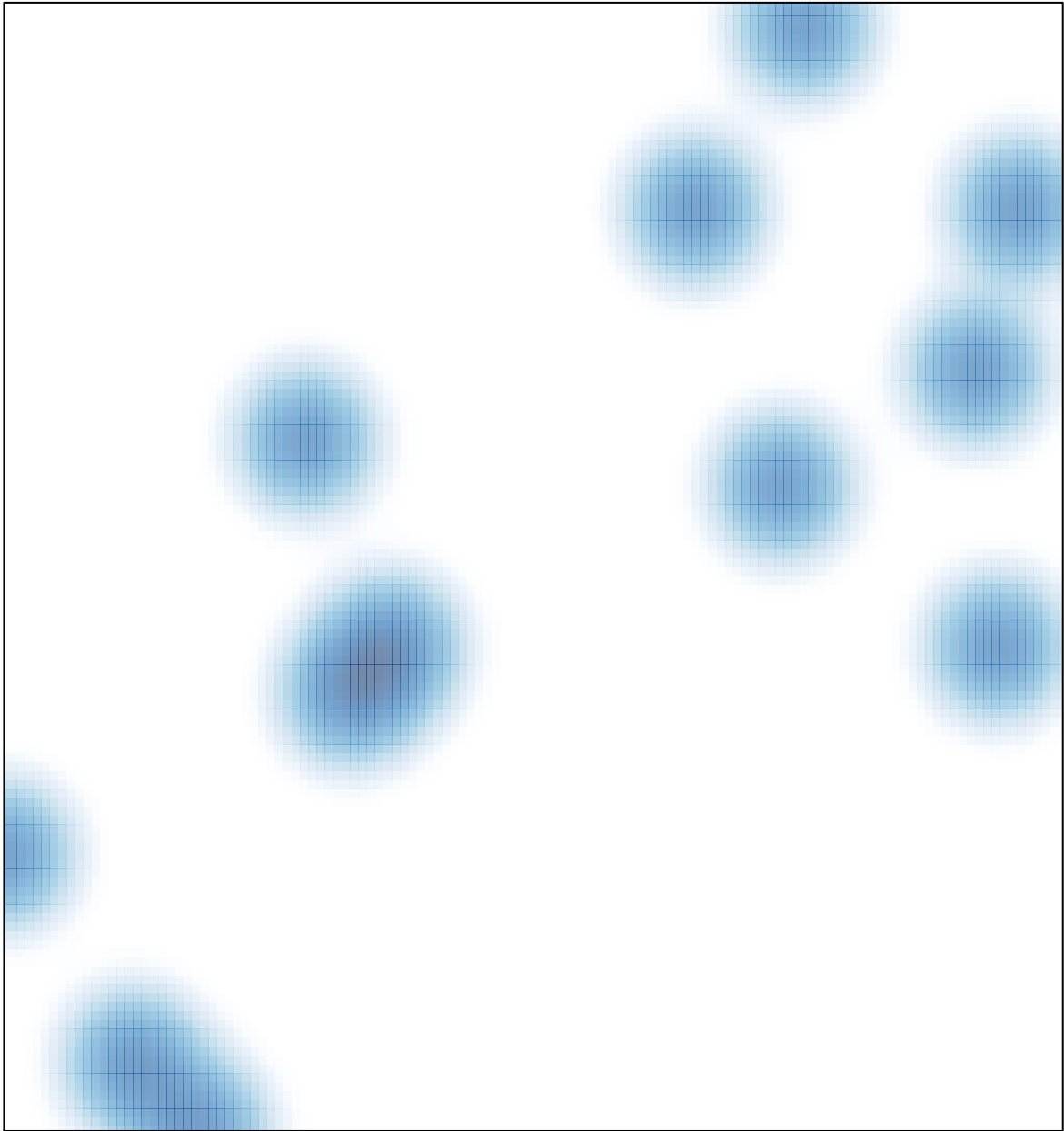


# MAGRANGEAS\_MULTIPLE\_MYELOMA\_IGG\_VS\_IGA\_DN



# features = 16  
chi-square p = 0.83

# AGRANGEAS\_MULTIPLE\_MYELOMA\_IGG\_VS\_IGA\_



# features = 16 , max = 1