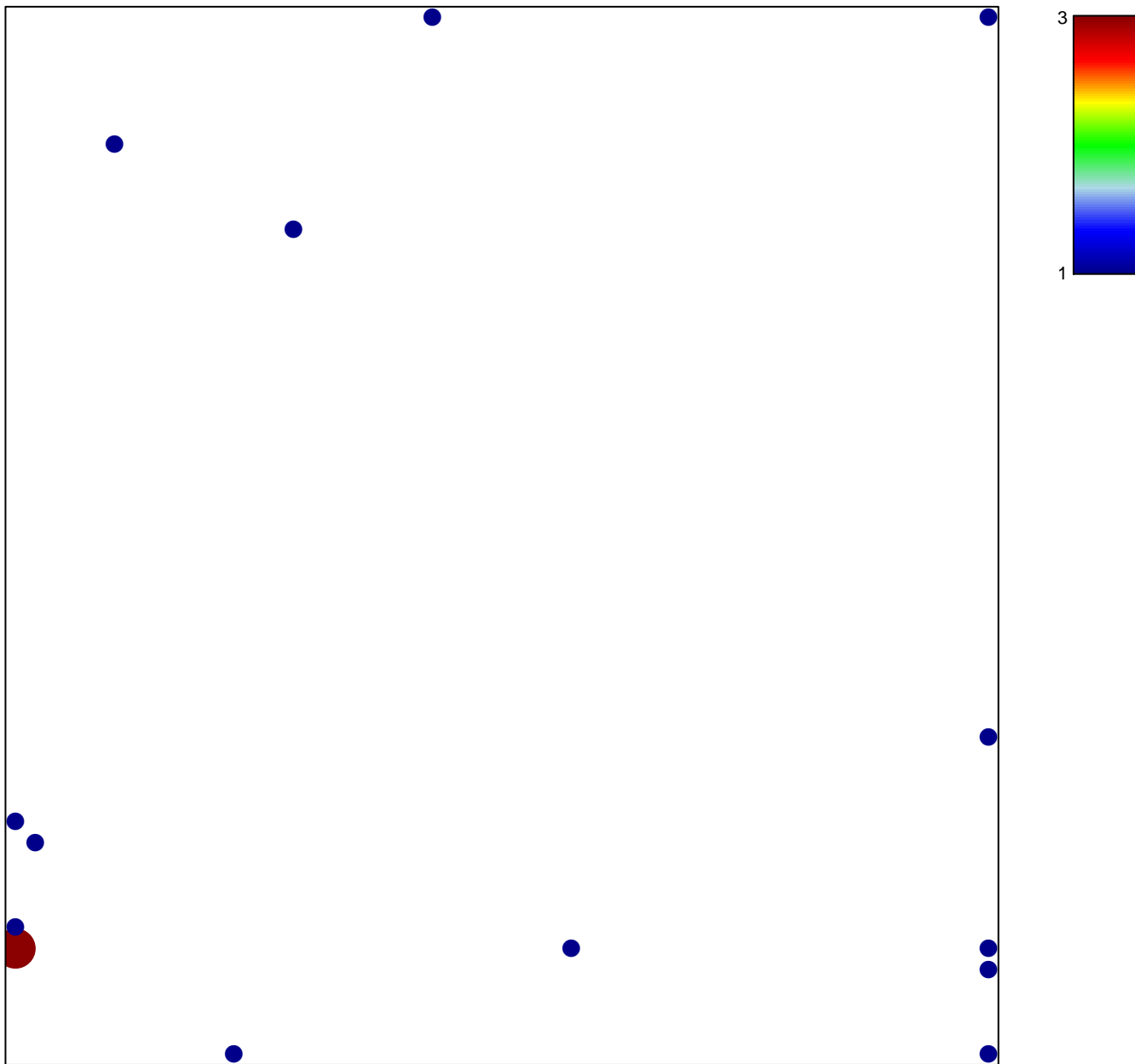
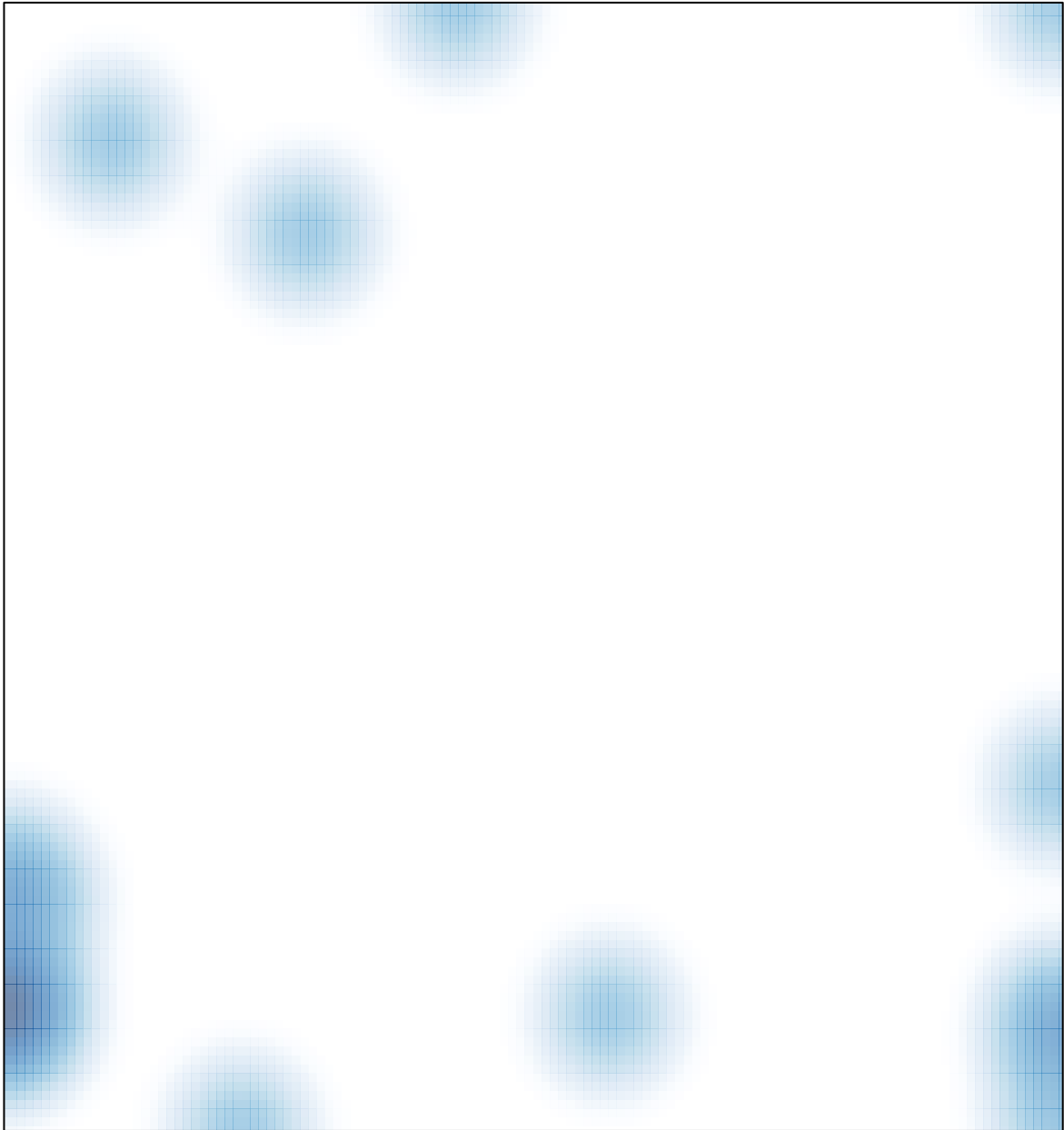


# ABRAHAM\_ALPC\_VS\_MULTIPLE\_MYELOMA\_UP



# features = 16  
chi-square p = 0.74

# ABRAHAM\_ALPC\_VS\_MULTIPLE\_MYELOMA\_UP



# features = 16 , max = 3