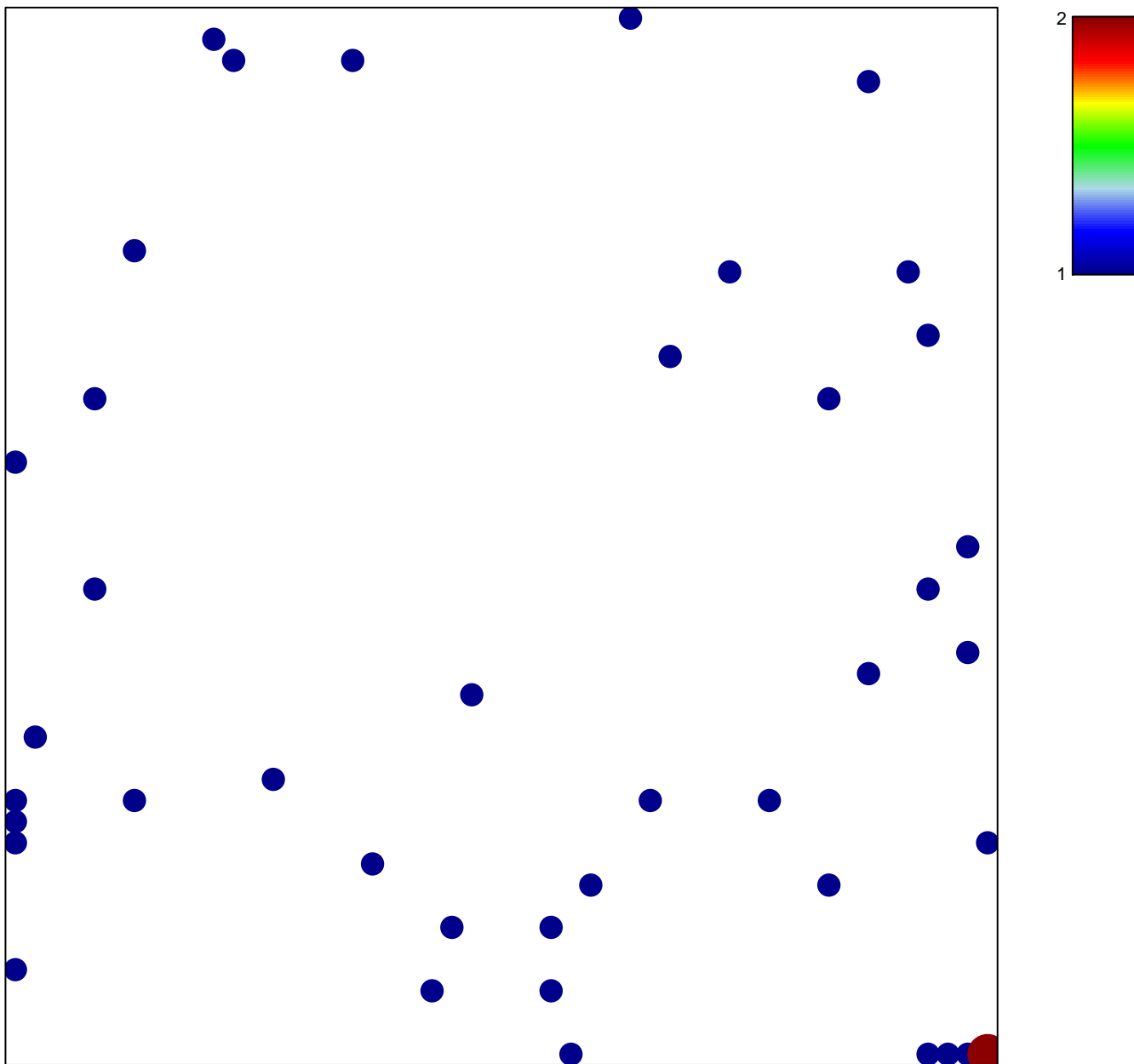
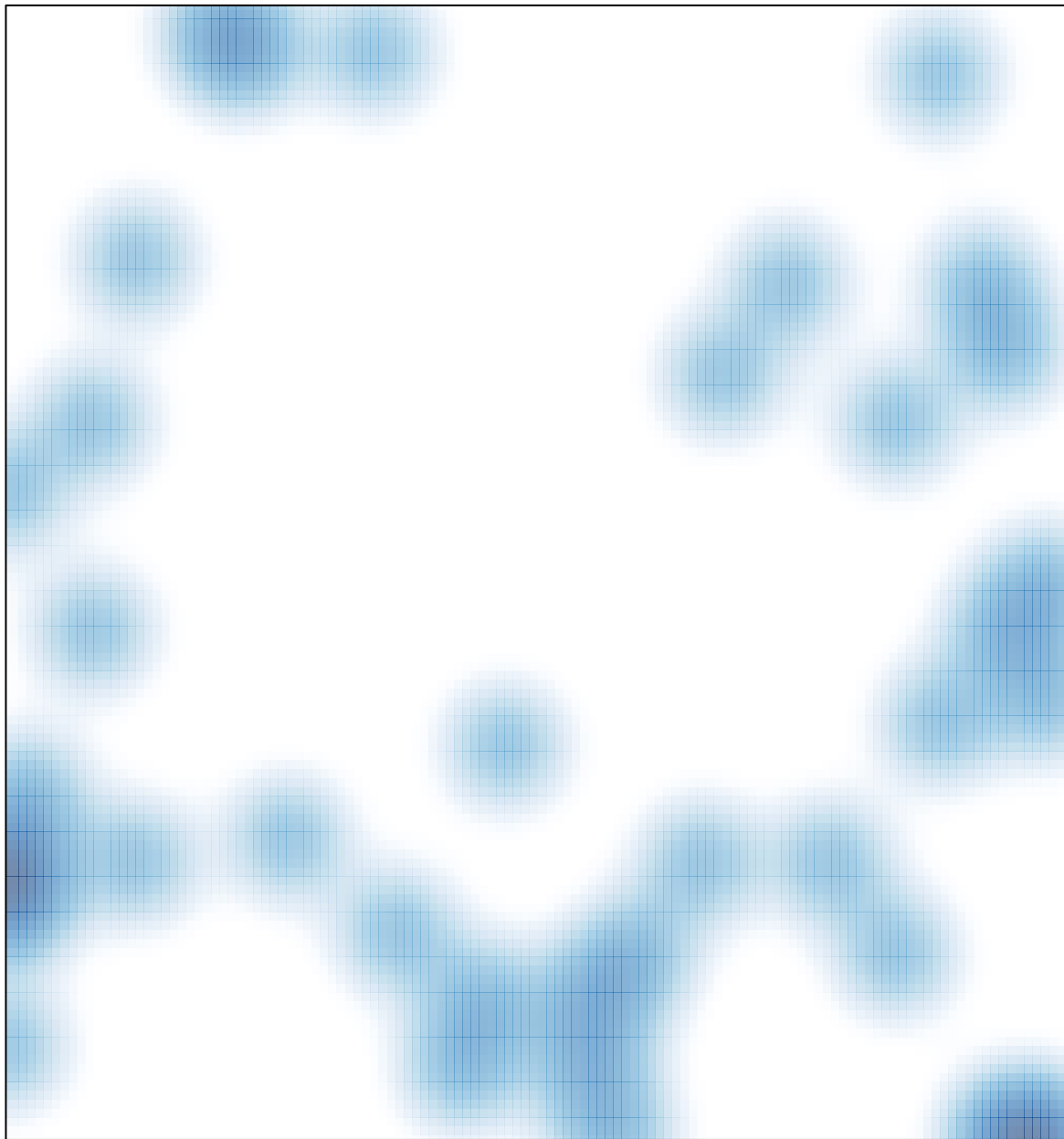


# ENGELMANN\_CANCER\_PROGENITORS\_UP



# features = 42  
chi-square p = 0.81

# ENGELMANN\_CANCER\_PROGENITORS\_UP



# features = 42 , max = 2