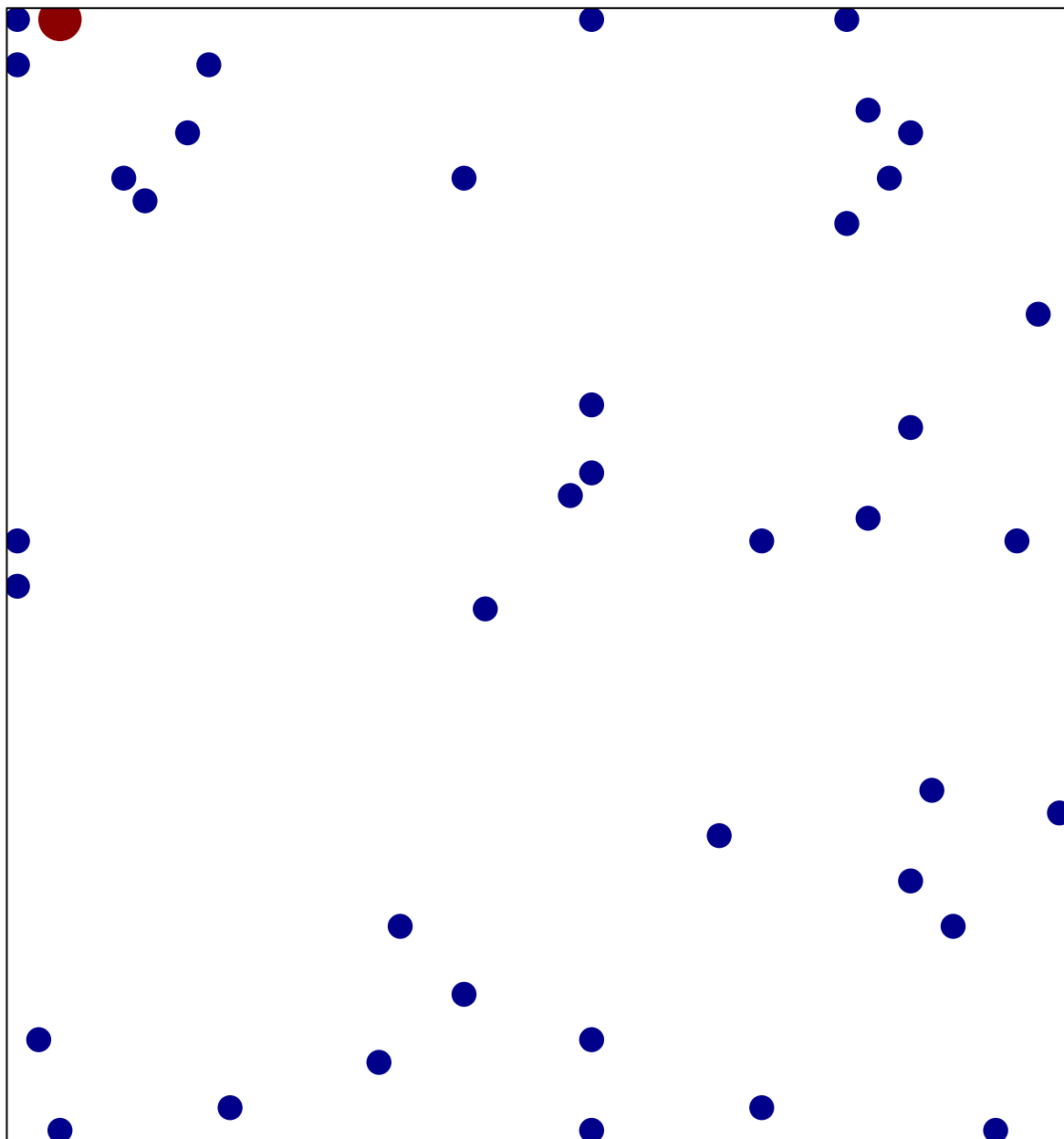
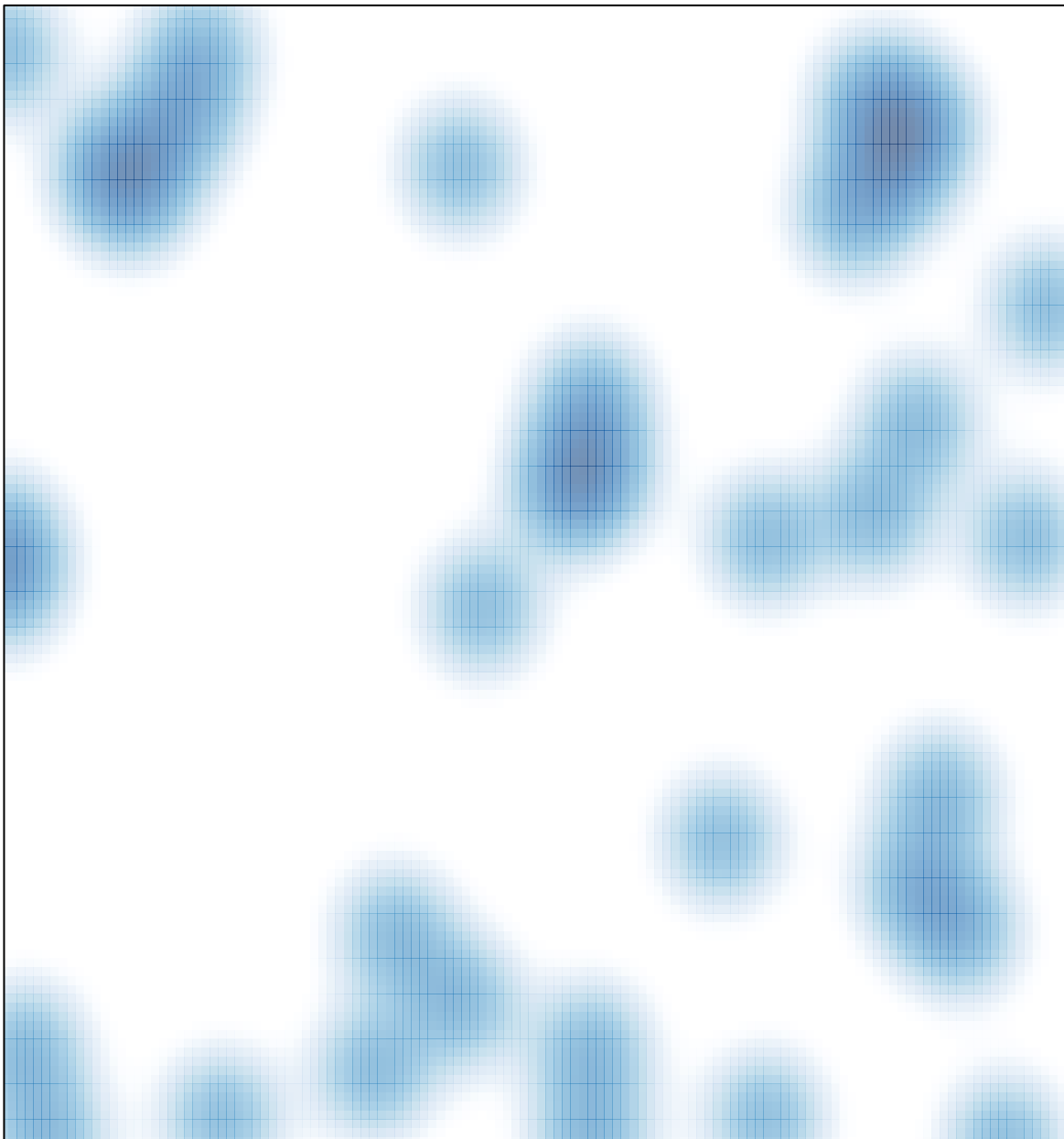


# BOYLAN\_MULTIPLE\_MYELOMA\_C\_UP



# features = 41  
chi-square p = 0.81

# BOYLAN\_MULTIPLE\_MYELOMA\_C\_UP



# features = 41 , max = 2