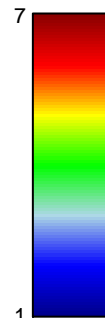
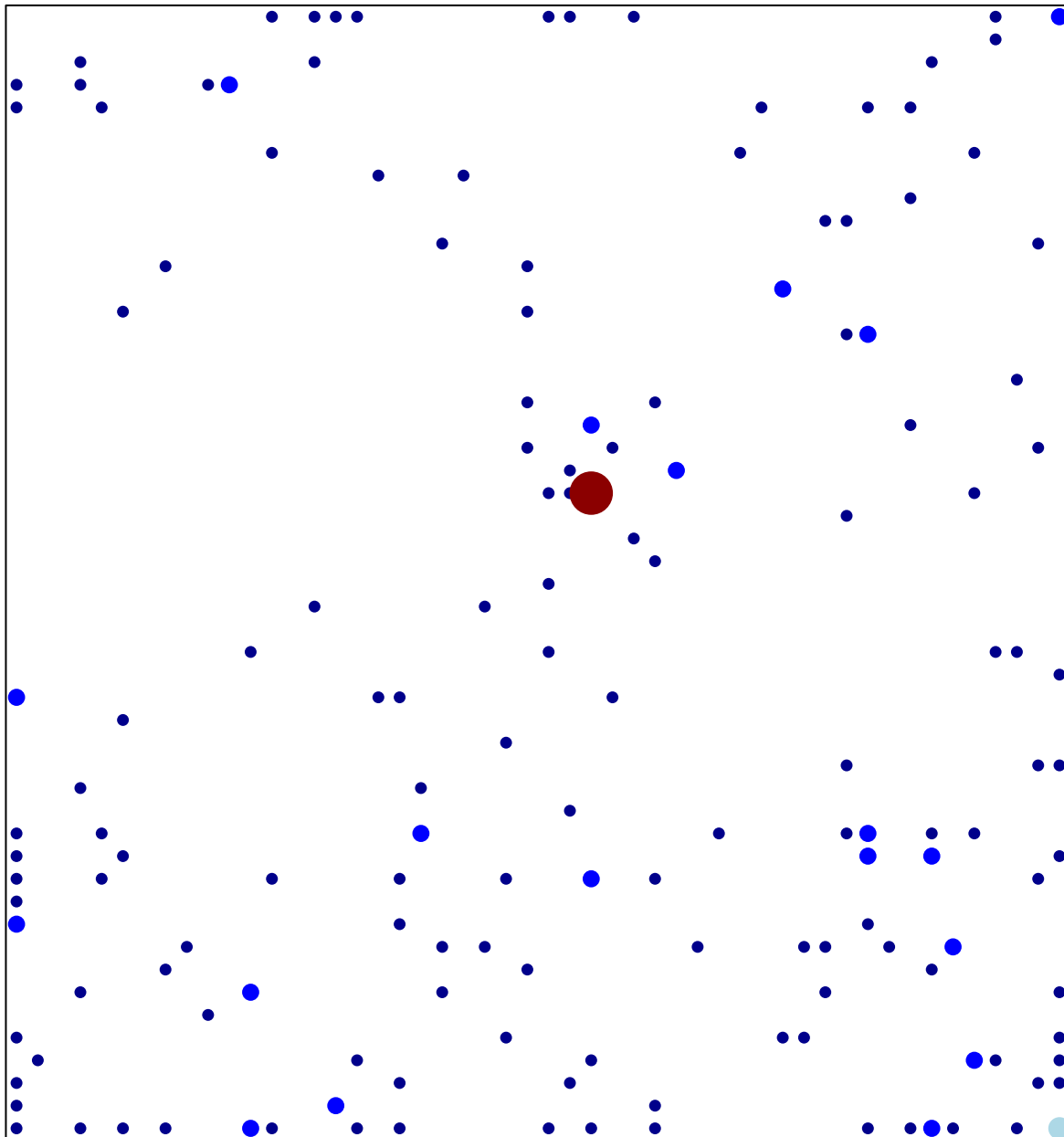
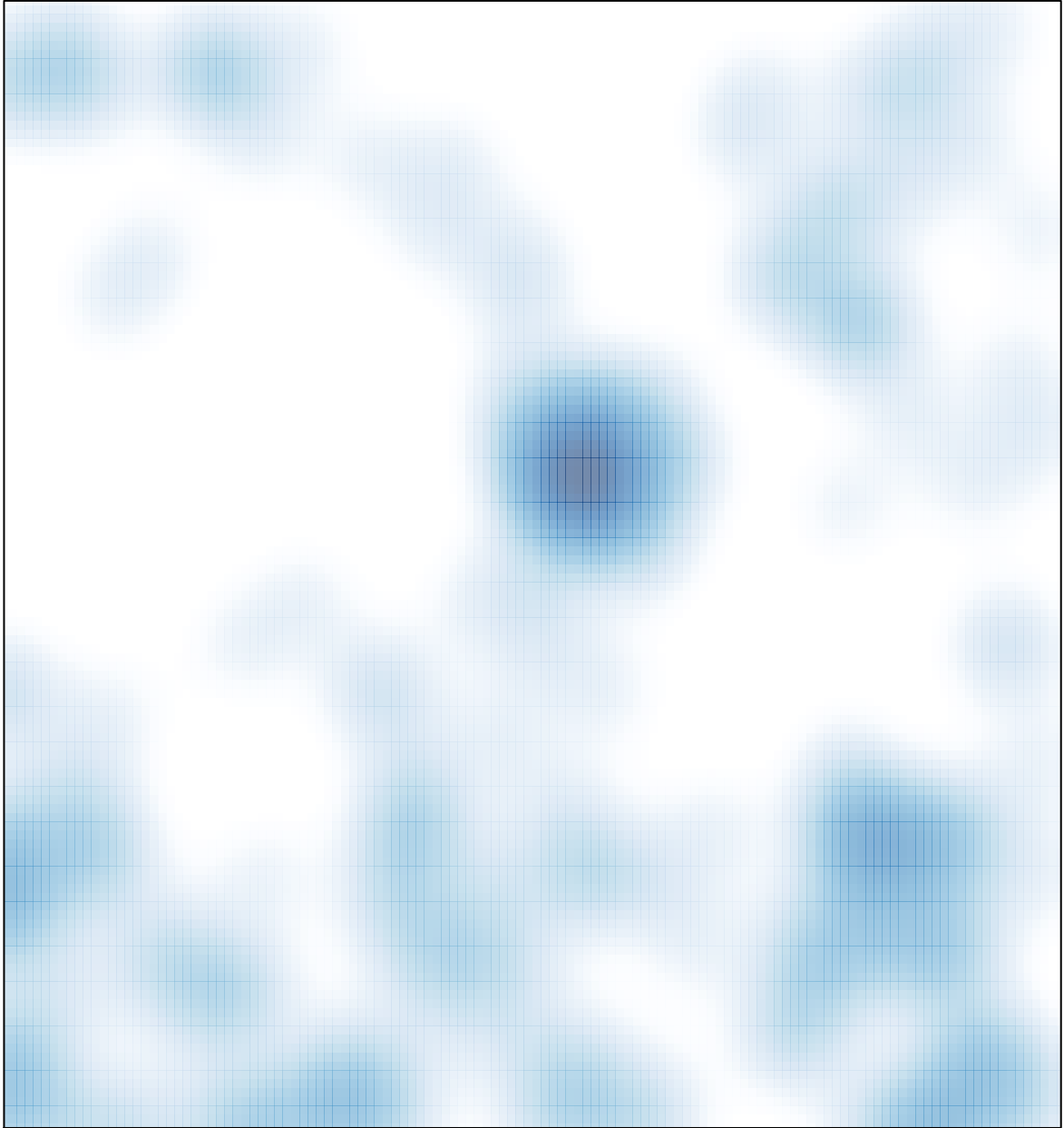


# BOYLAN\_MULTIPLE\_MYELOMA\_C\_D\_DN



# features = 181  
chi-square p = 0.72

# BOYLAN\_MULTIPLE\_MYELOMA\_C\_D\_DN



# features = 181 , max = 7