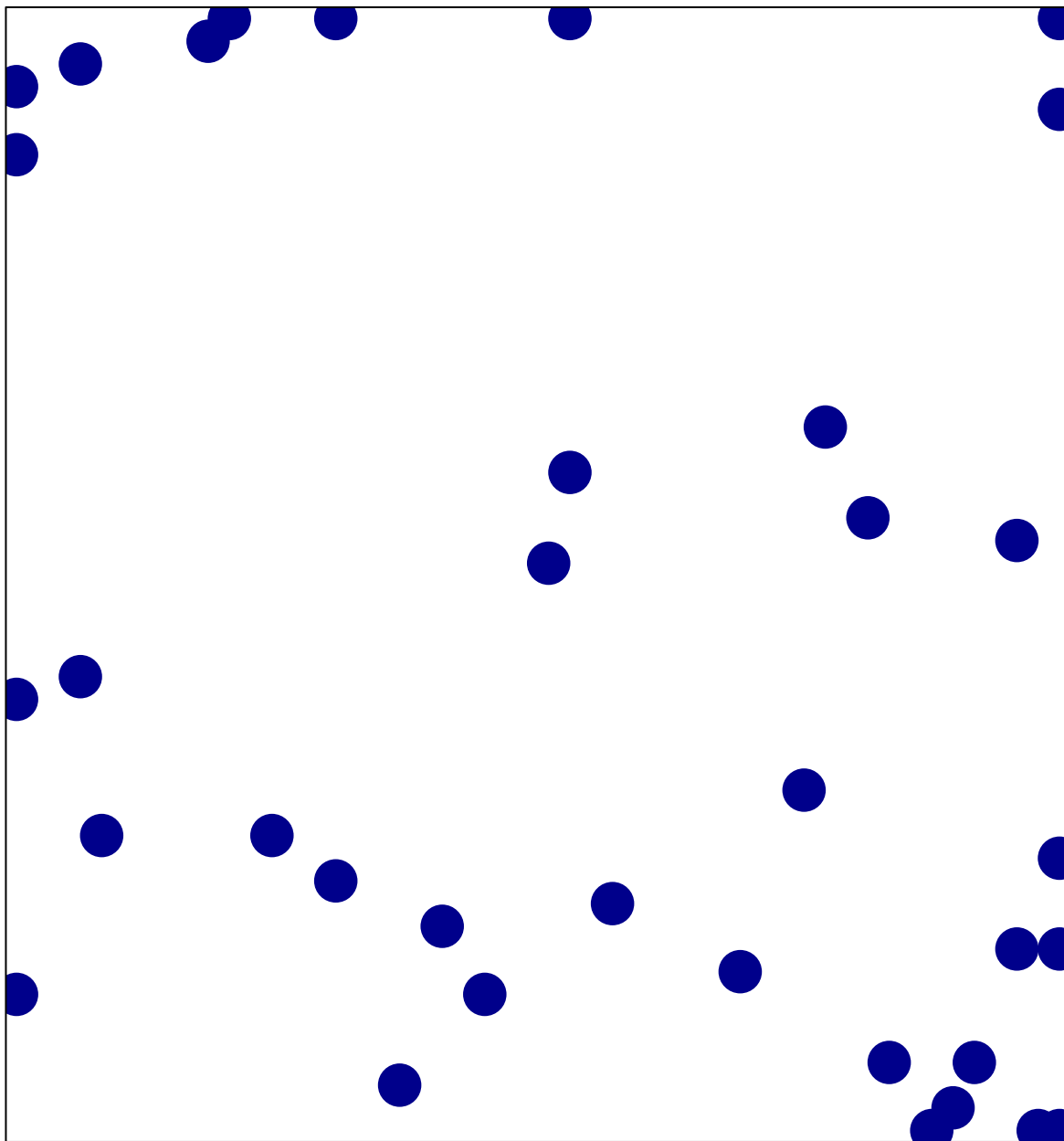
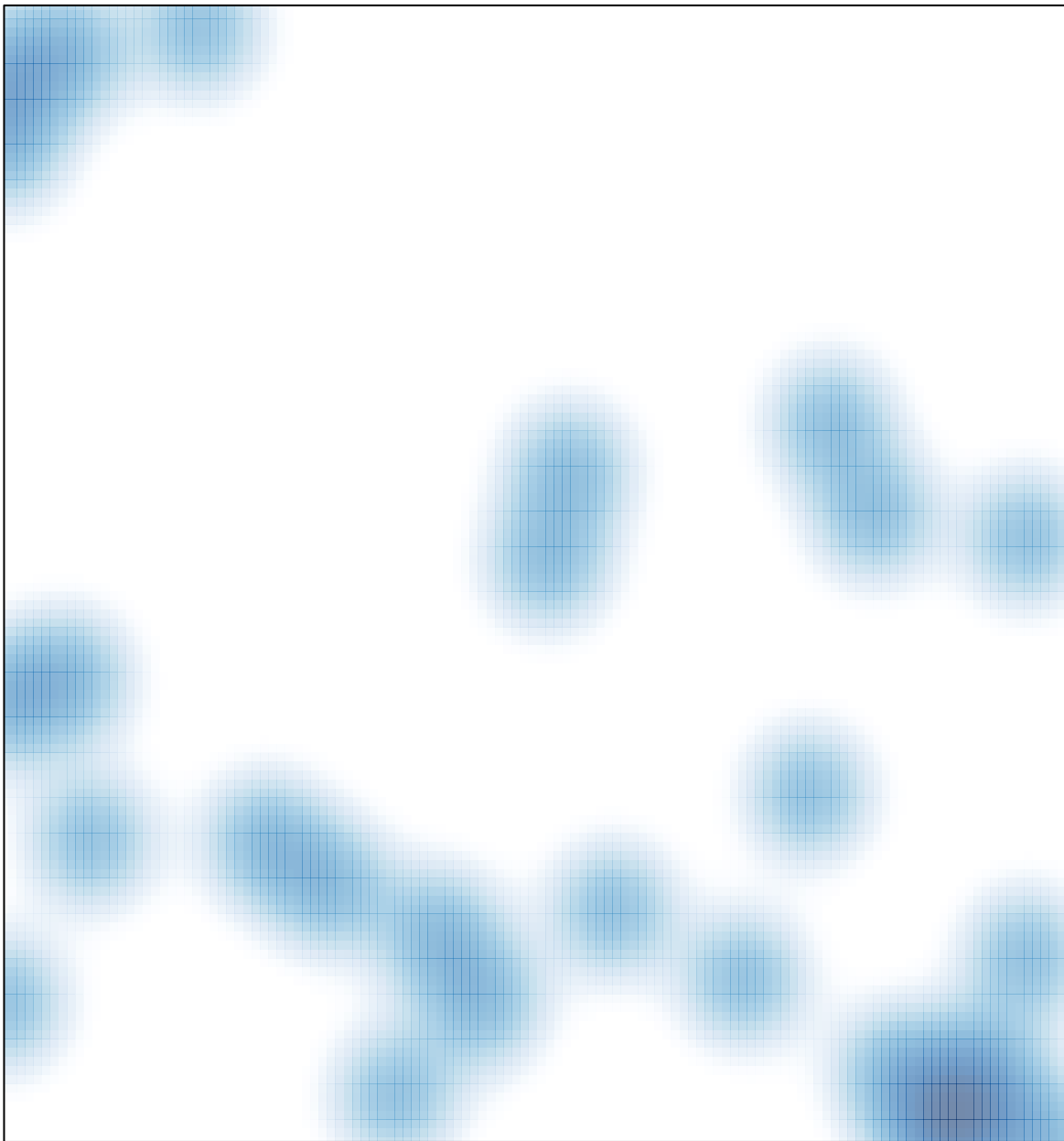


# BOYLAN\_MULTIPLE\_MYELOMA\_D\_CLUSTER\_DN



# features = 35  
chi-square p = 0.82

# BOYLAN\_MULTIPLE\_MYELOMA\_D\_CLUSTER\_DN



# features = 35 , max = 1