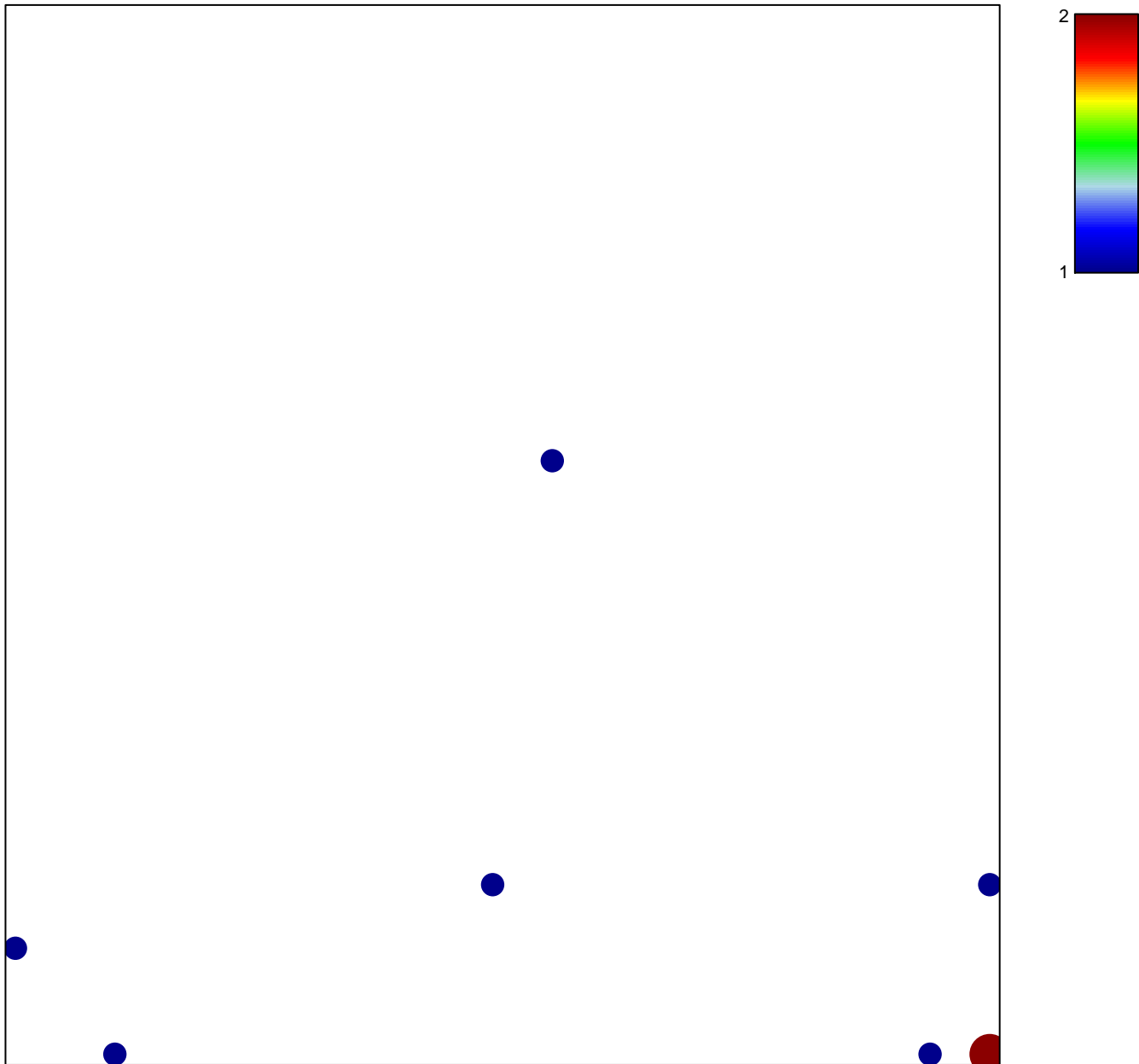
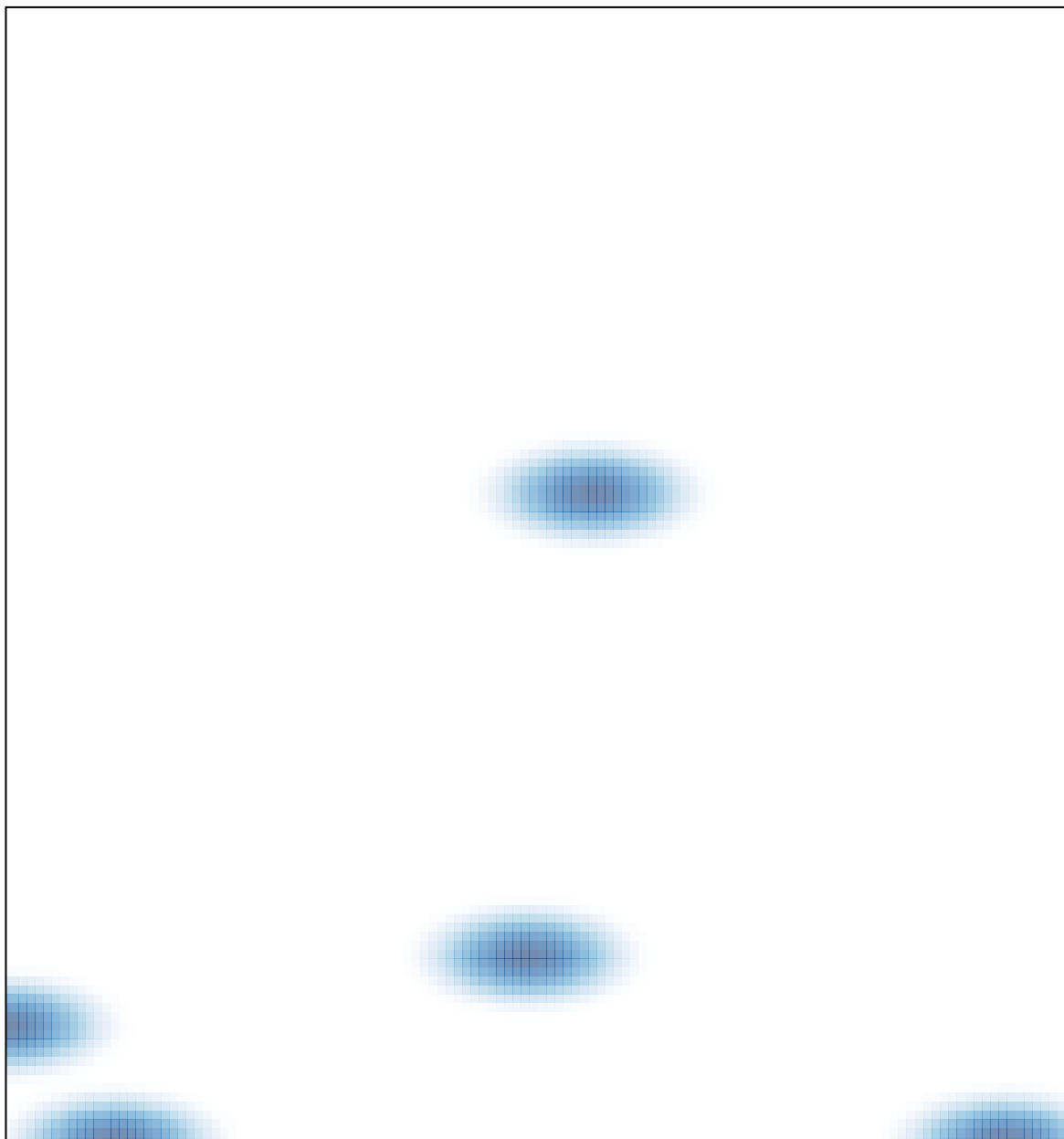


# BOYLAN\_MULTIPLE\_MYELOMA\_PCA1\_DN



# features = 8  
chi-square p = 0.77

# BOYLAN\_MULTIPLE\_MYELOMA\_PCA1\_DN



# features = 8 , max = 2