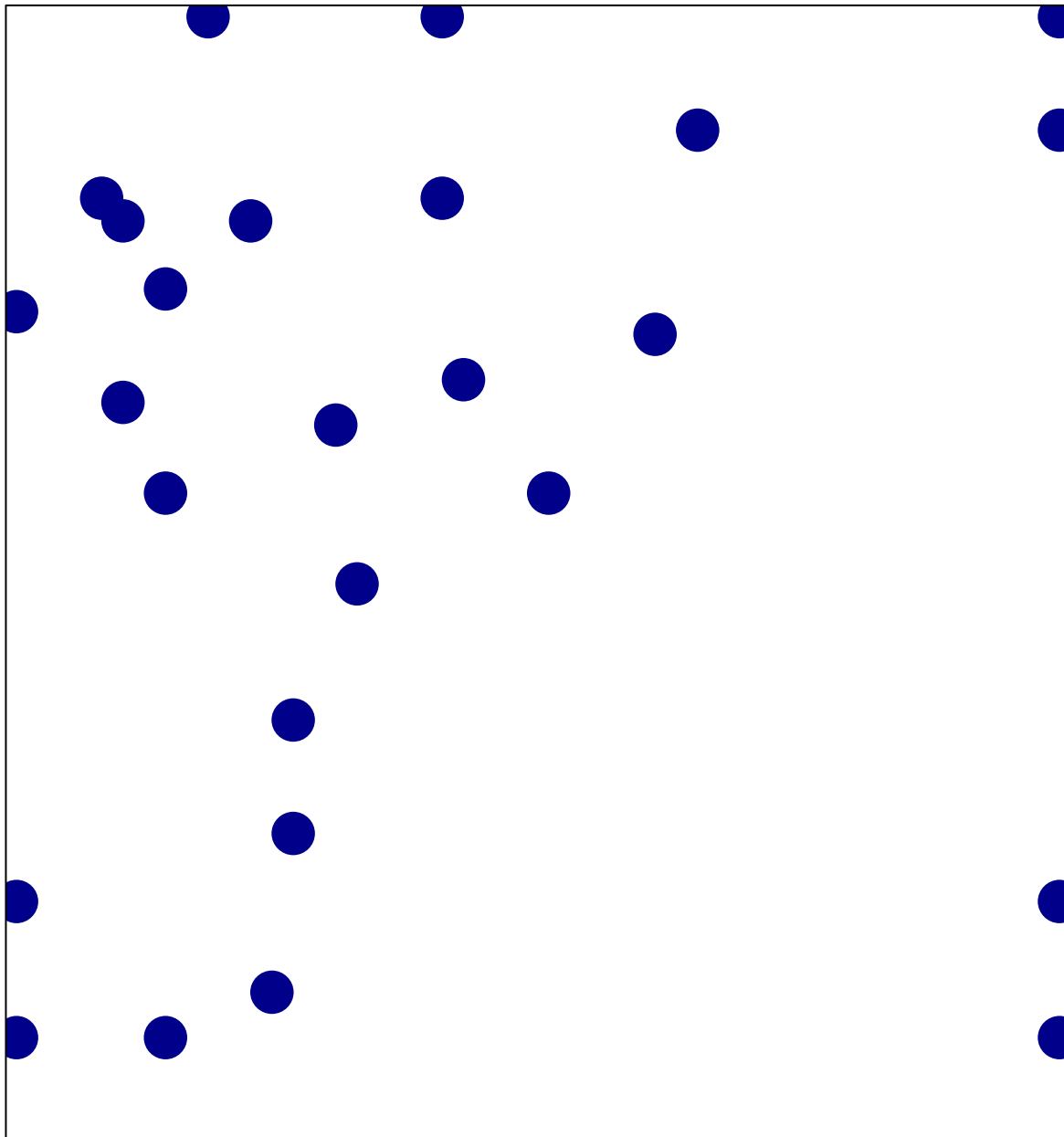


# BARRIER\_CANCER\_RELAPSE\_NORMAL\_SAMPLE\_UP



# features = 26  
chi-square p = 0.81

BARRIER\_CANCER\_RELAPSE\_NORMAL\_SAMPLE\_U



# features = 26 , max = 1