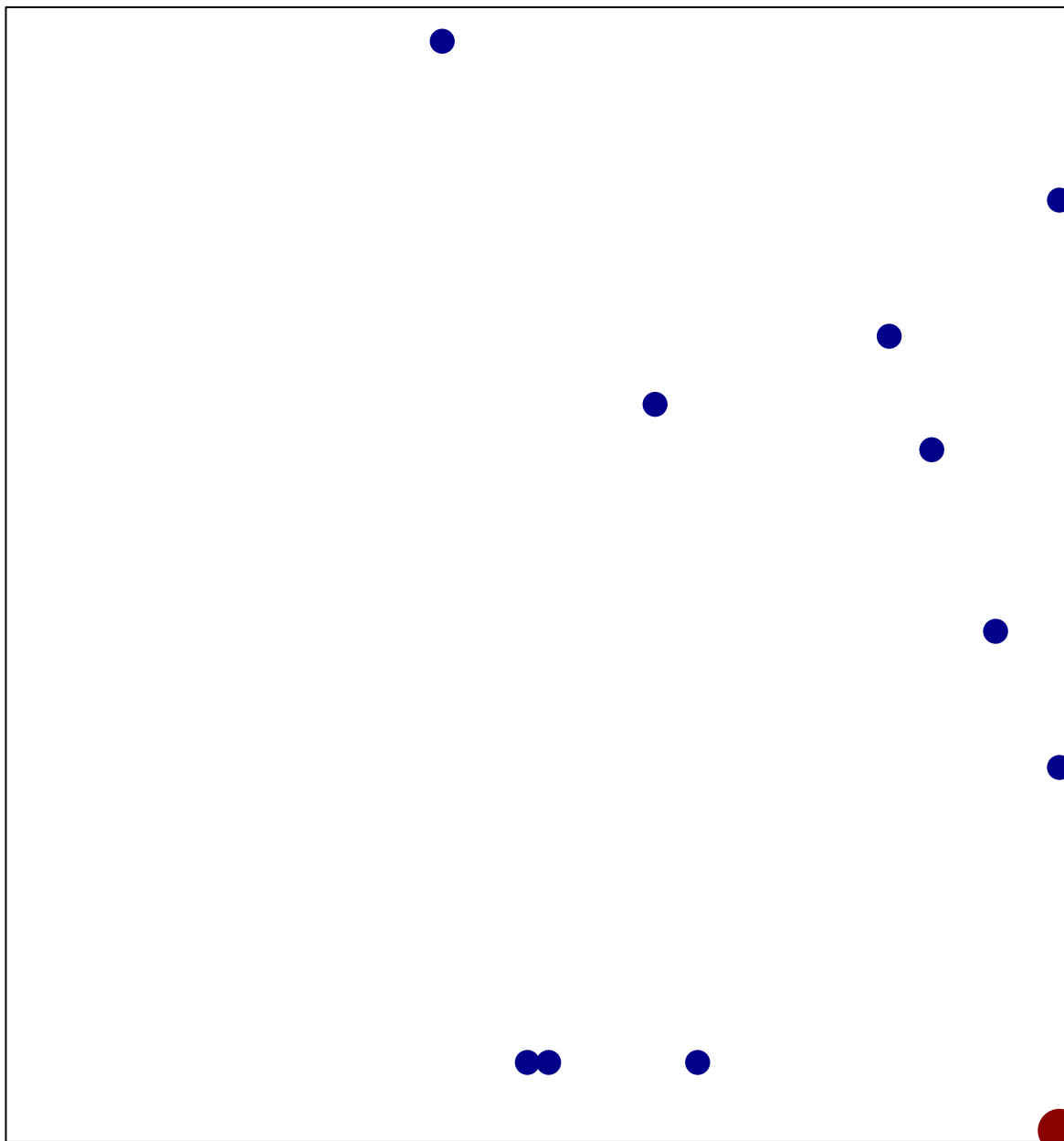


# NIELSEN\_MALIGNAT\_FIBROUS\_HISTIOCYTOMA\_UP



# features = 12  
chi-square p = 0.8

# NIELSEN\_MALIGNANT\_FIBROUS\_HISTIOCYTOMA\_U



# features = 12 , max = 2