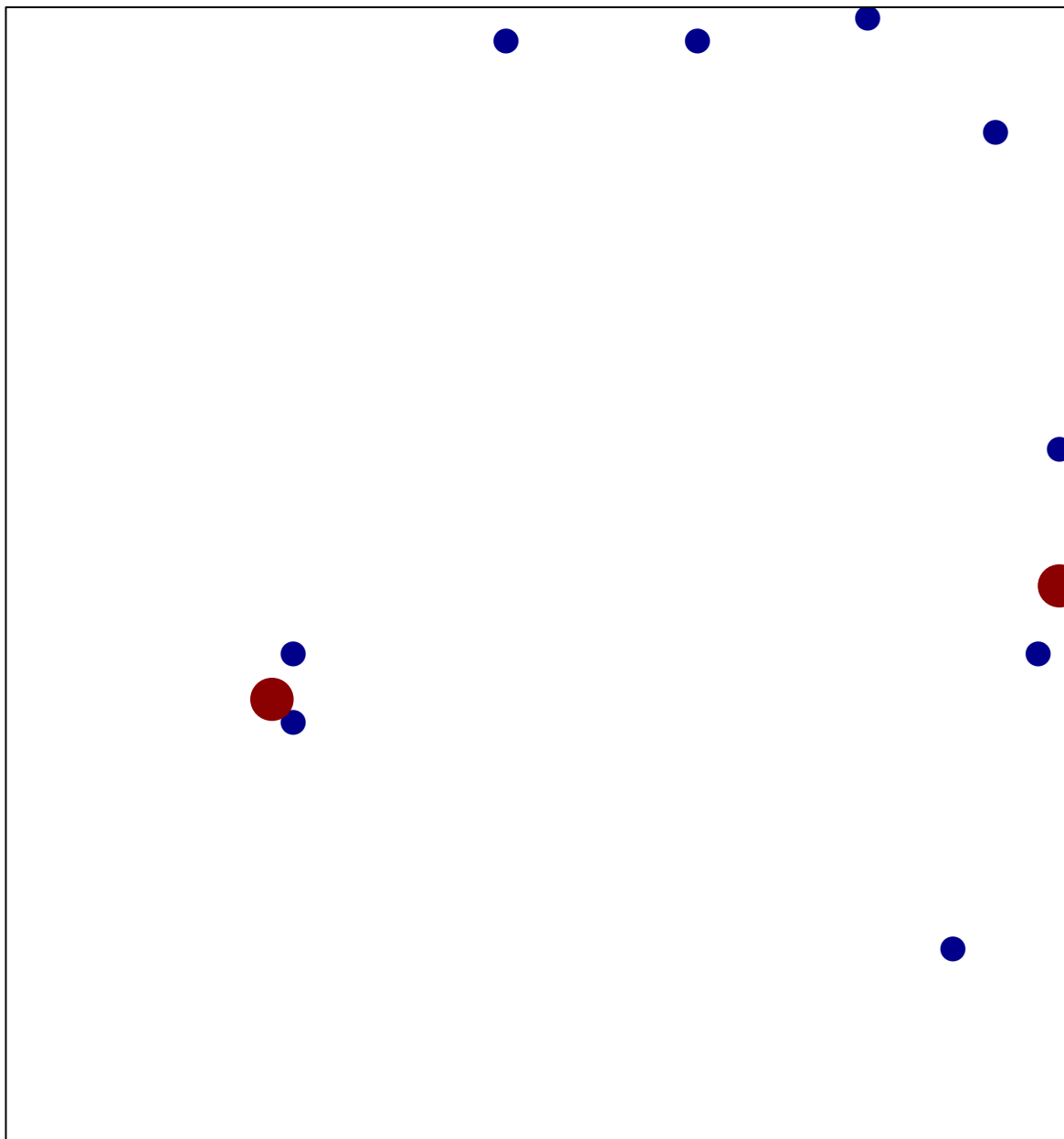


CHNG_MULTIPLE_MYELOMA_HYPERPLOID_UP



features = 13
chi-square p = 0.81

CHNG_MULTIPLE_MYELOMA_HYPERPLOID_UP



features = 13 , max = 2