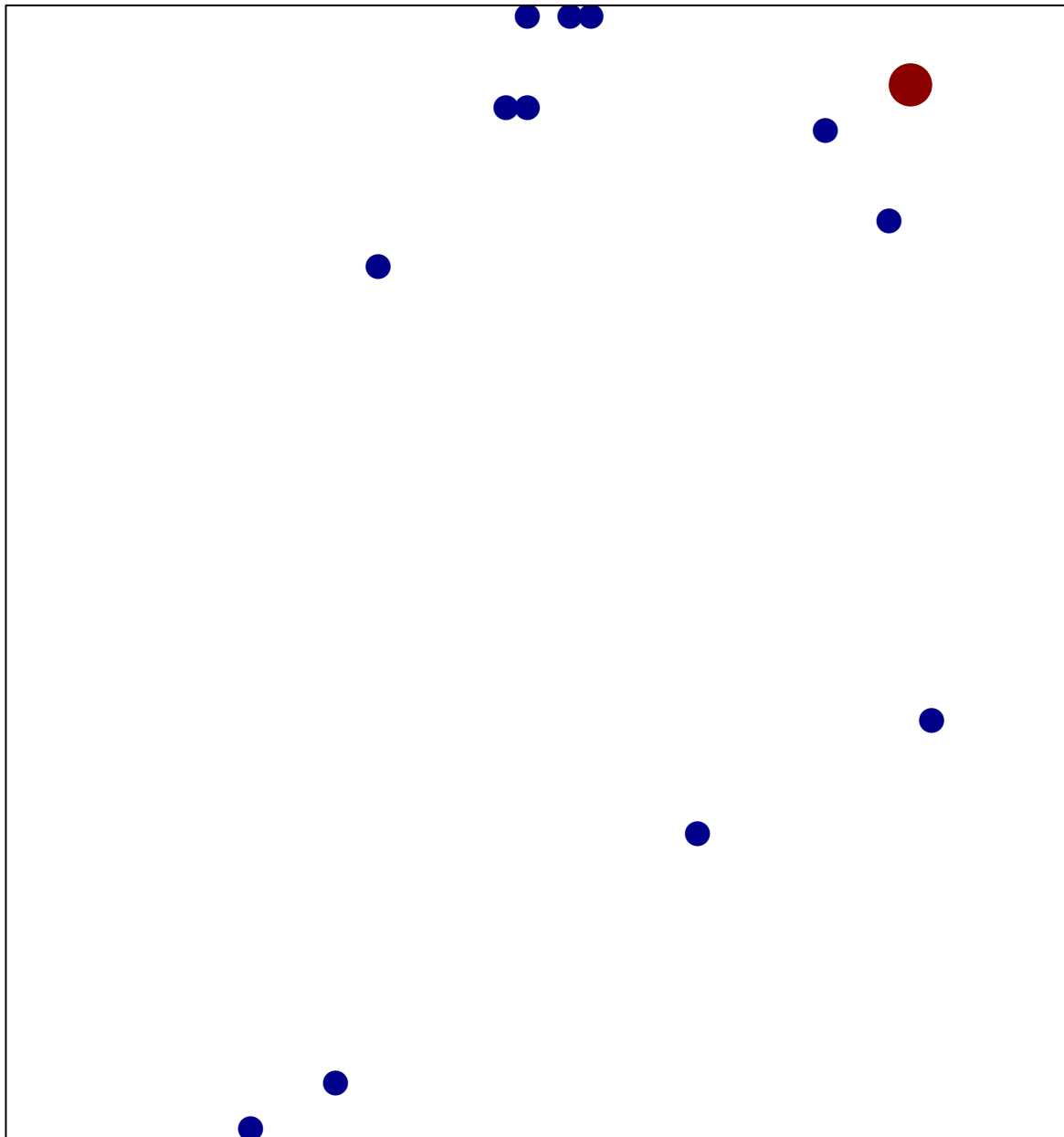


CHNG_MULTIPLE_MYELOMA_HYPERPLOID_DN



features = 14
chi-square p = 0.83

CHNG_MULTIPLE_MYELOMA_HYPERPLOID_DN



features = 14 , max = 2