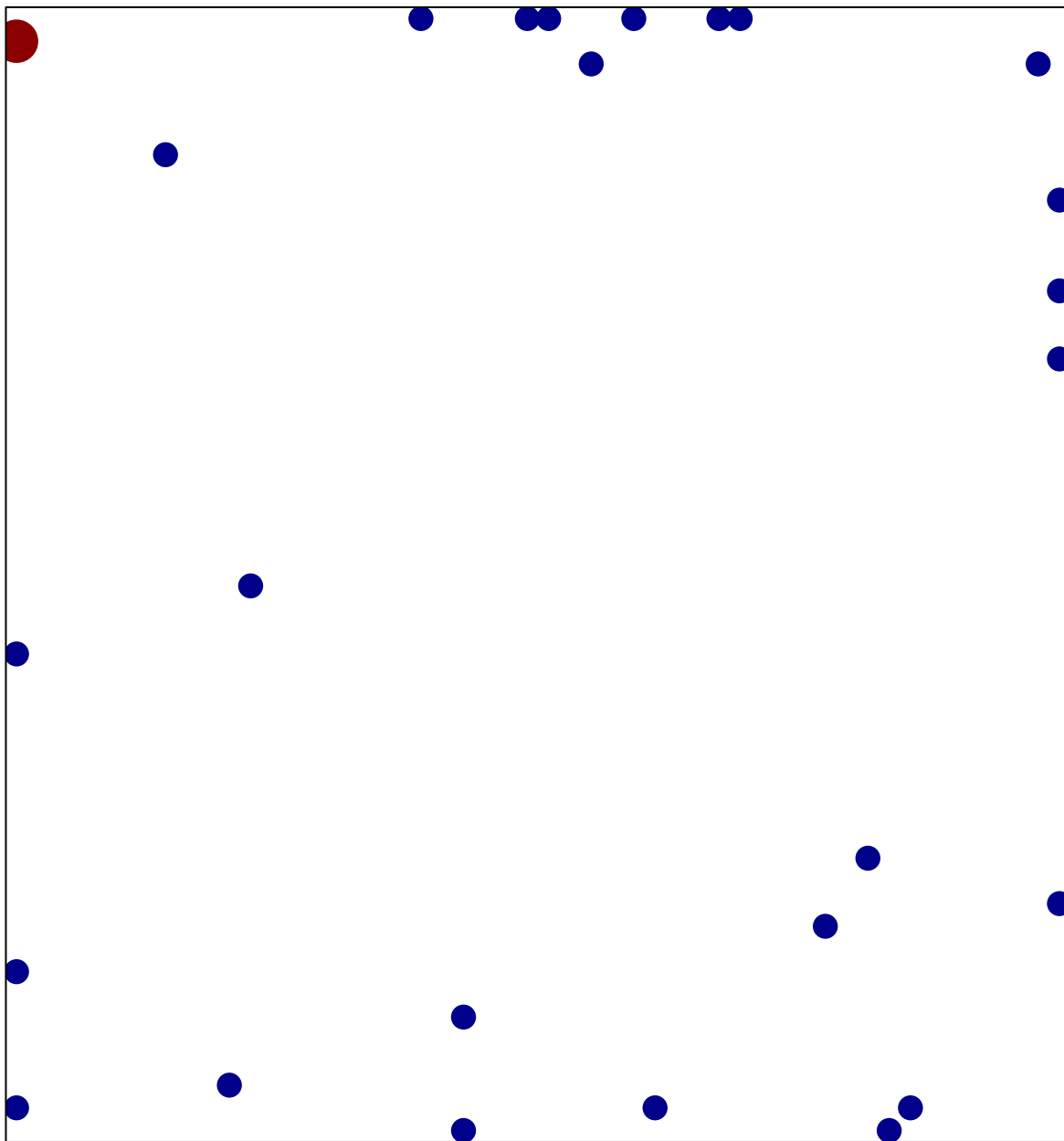
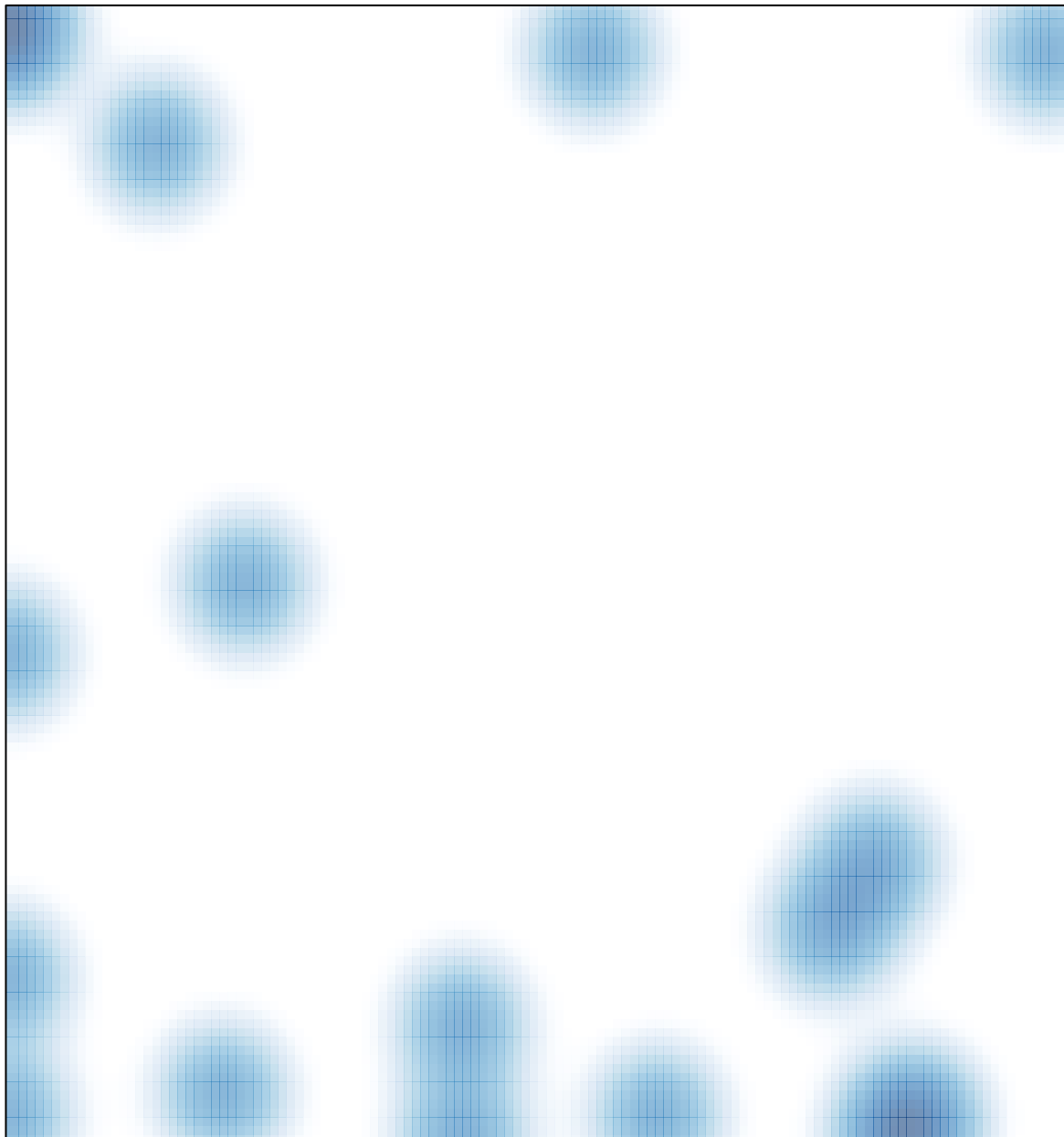


CHNG_MULTIPLE_MYELOMA_HYPERPLOID_DN



features = 27
chi-square p = 0.78

CHNG_MULTIPLE_MYELOMA_HYPERPLOID_DN



features = 27 , max = 2