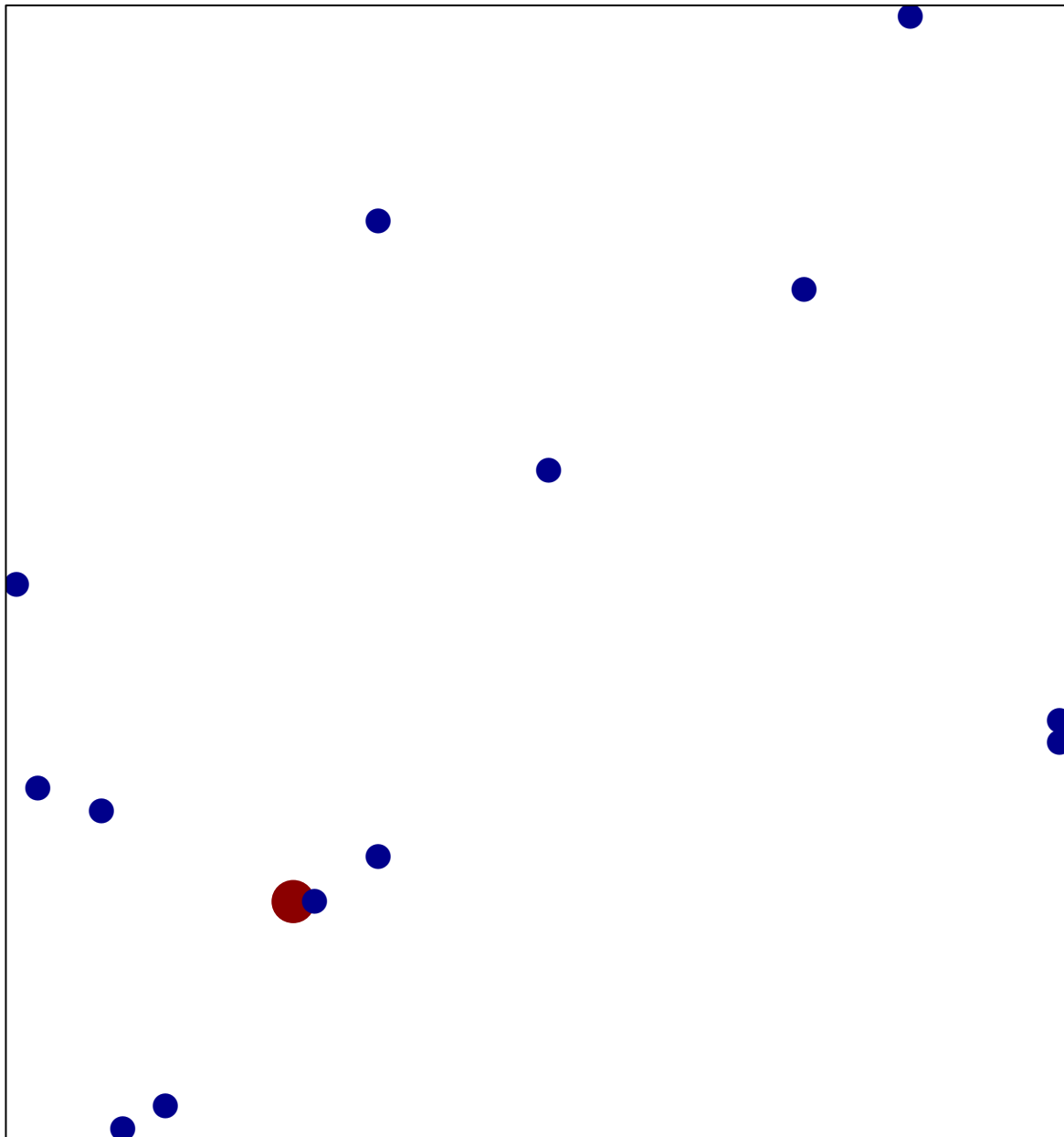
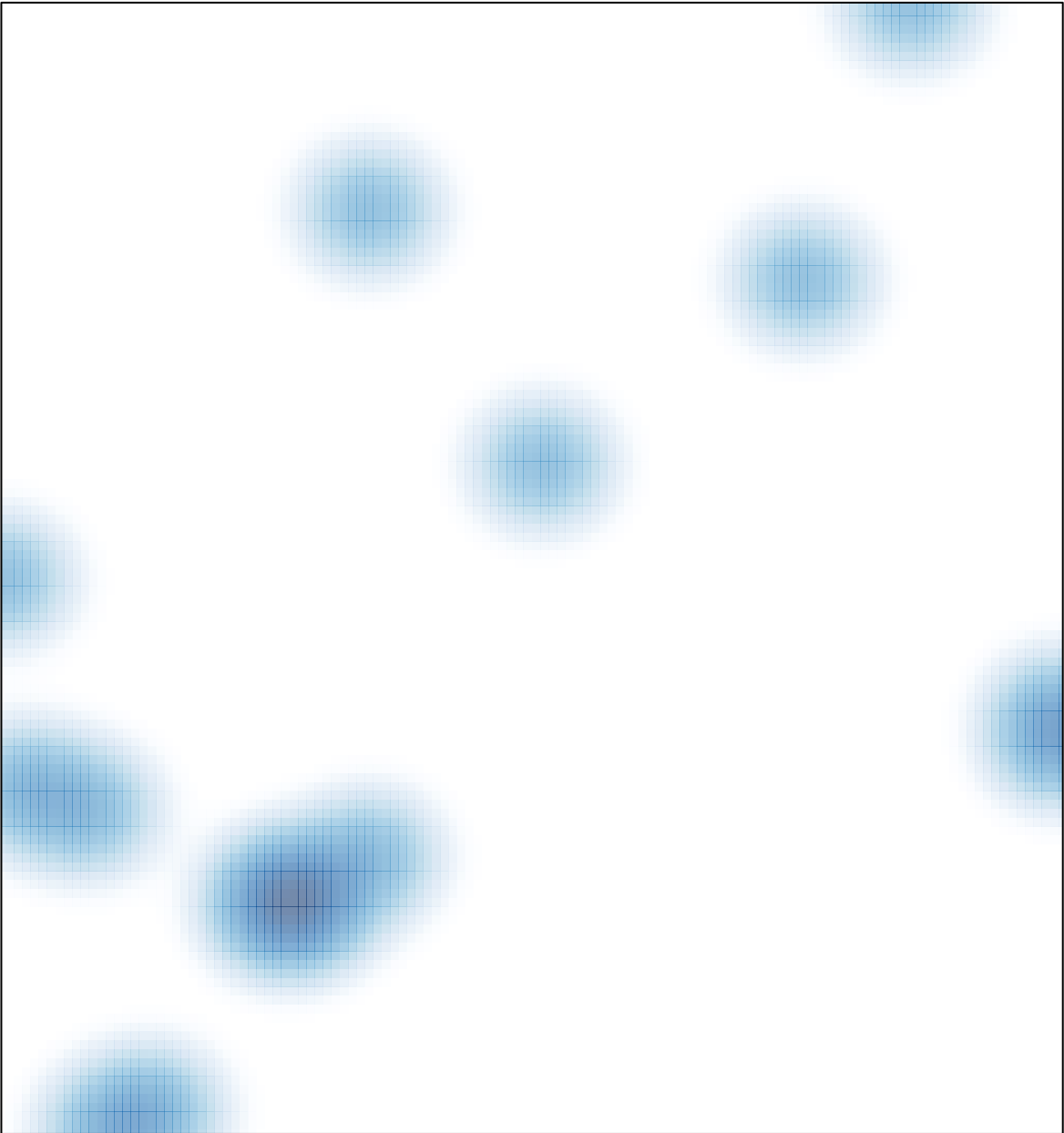


MATTIOLI_MULTIPLE_MYELOMA_SUBGROUPS



features = 15
chi-square p = 0.83

MATTIOLI_MULTIPLE_MYELOMA_SUBGROUPS



features = 15 , max = 2