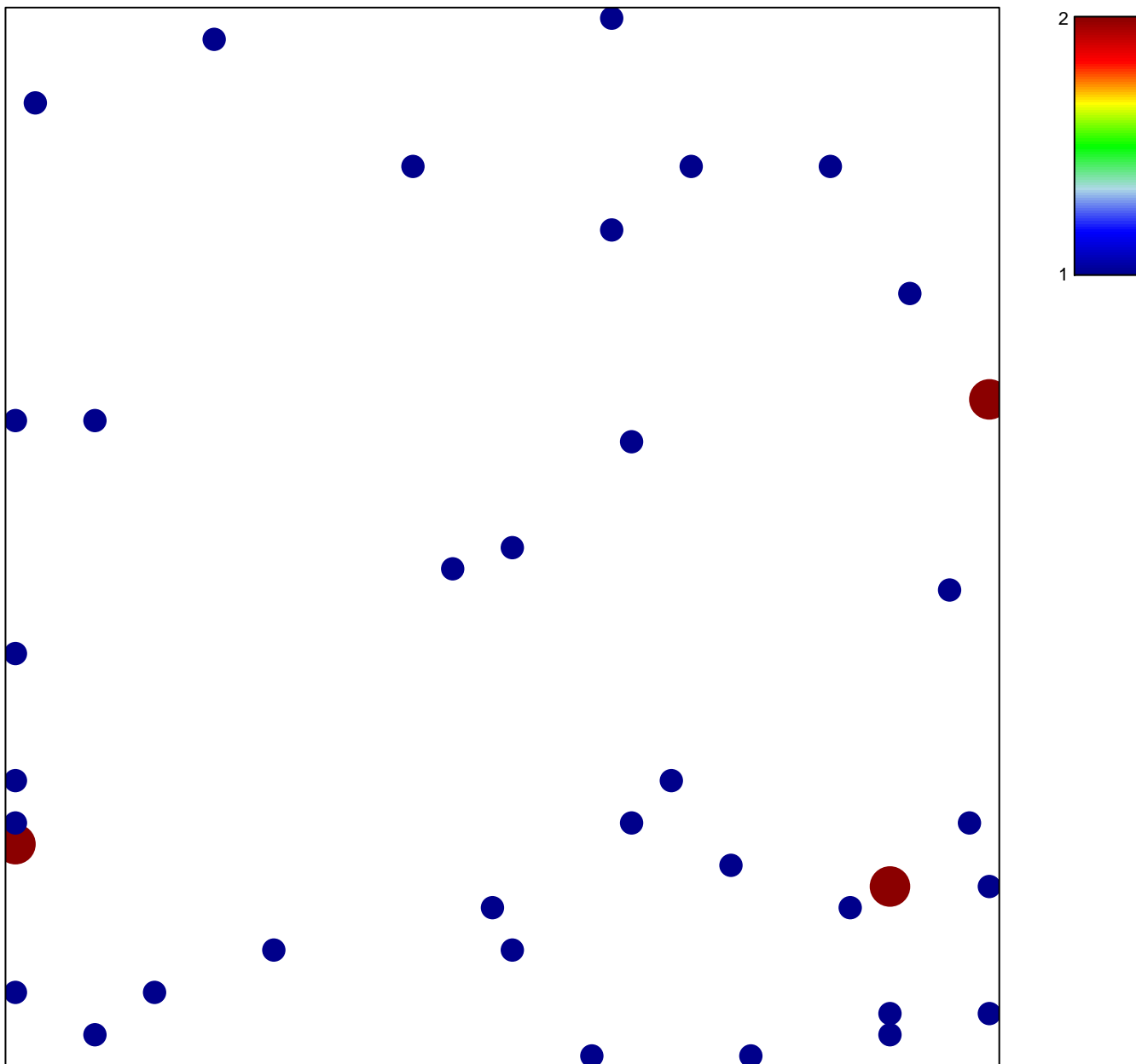
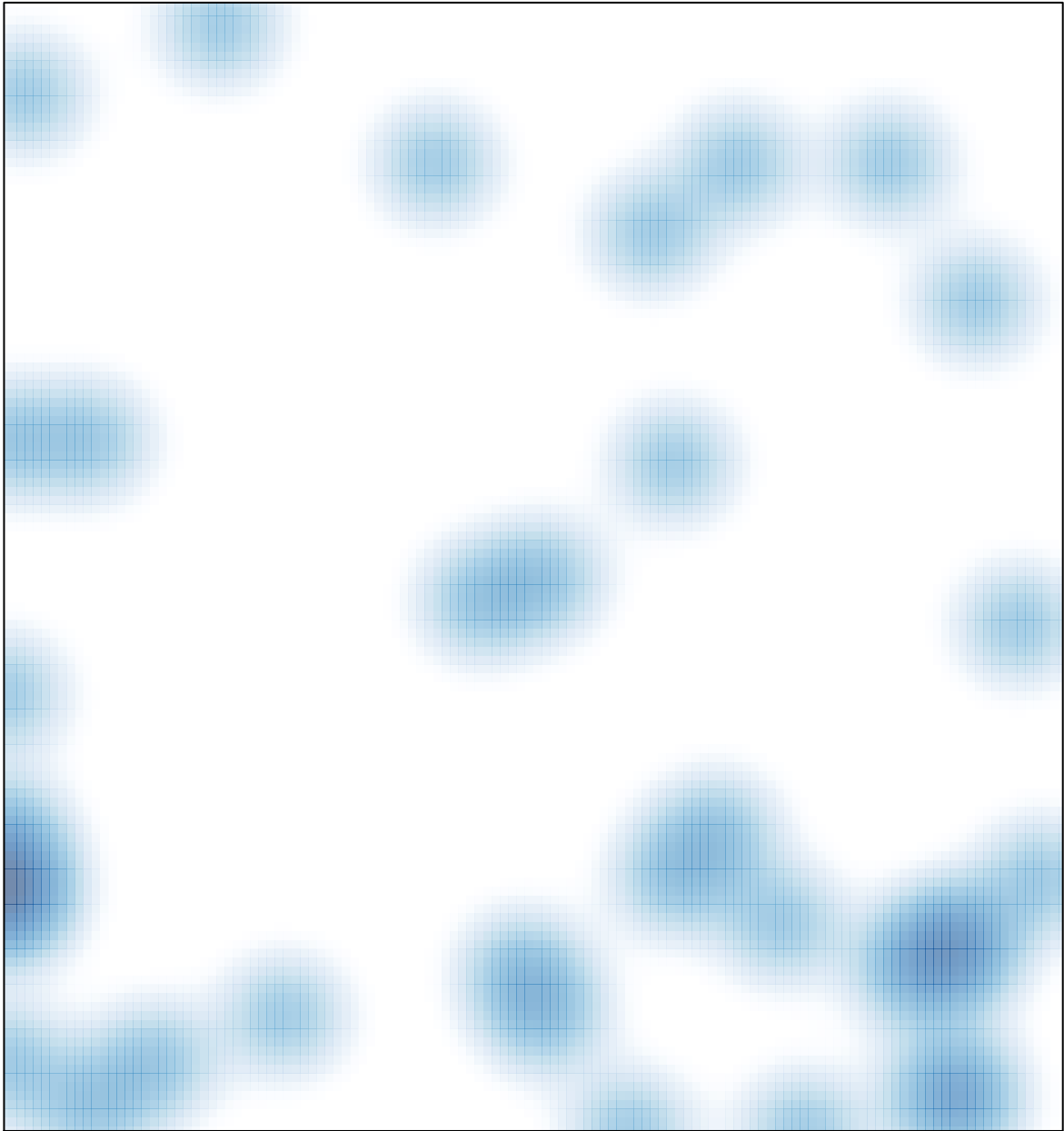


ZHAN_MULTIPLE_MYELOMA_CD1_DN



features = 40
chi-square p = 0.8

ZHAN_MULTIPLE_MYELOMA_CD1_DN



features = 40 , max = 2