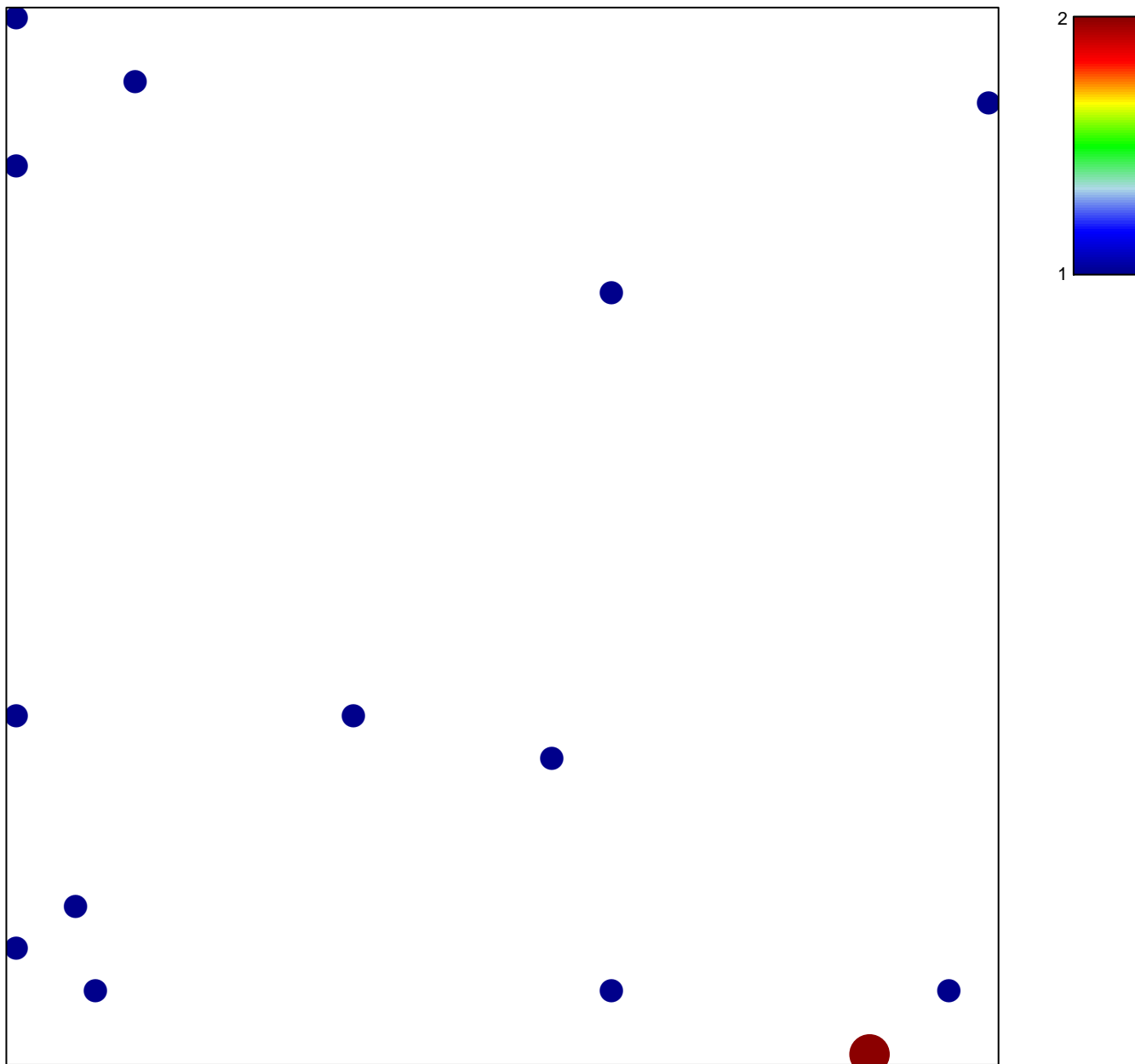
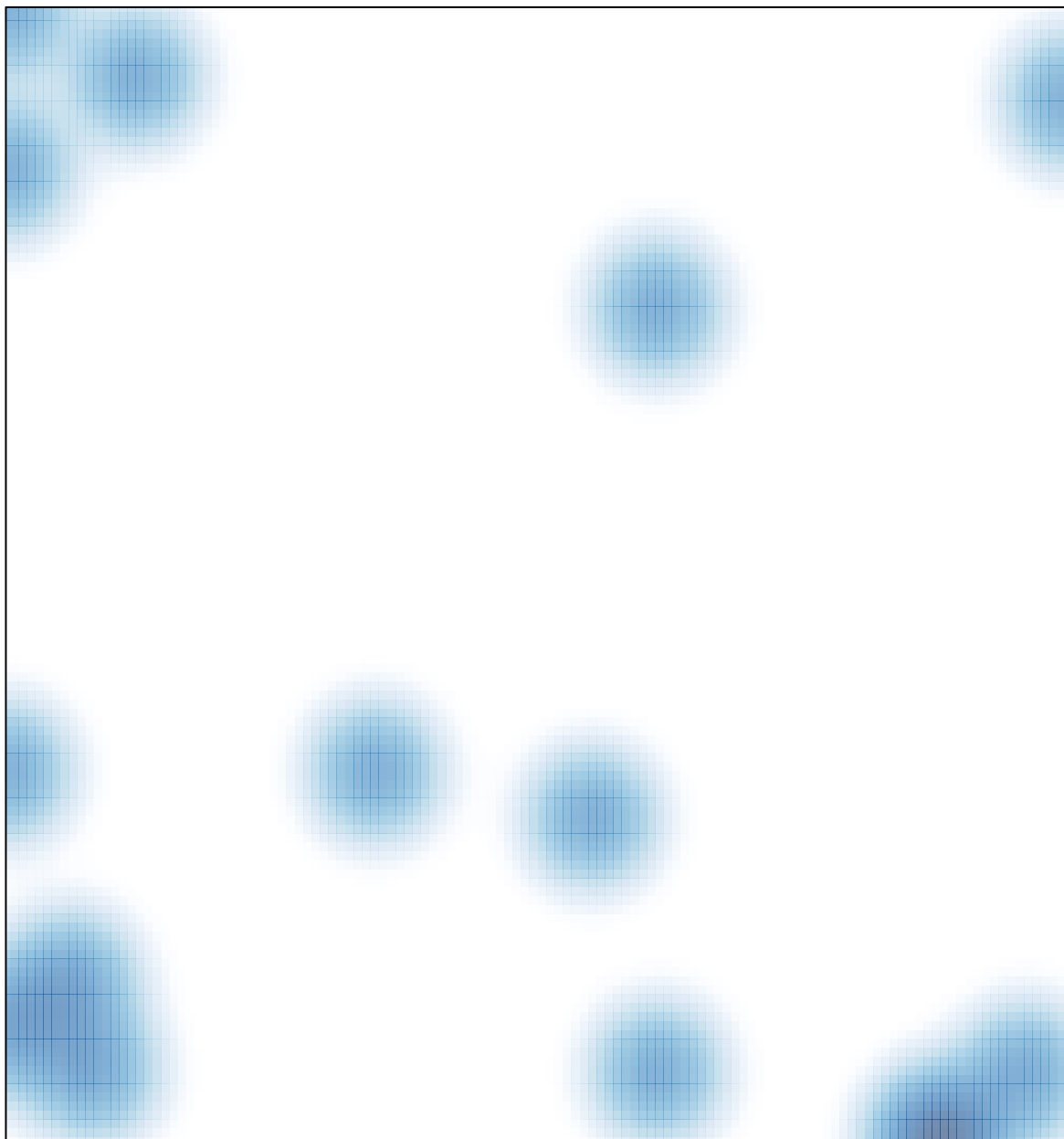


ZHAN_MULTIPLE_MYELOMA_CD1_UP



features = 15
chi-square p = 0.8

ZHAN_MULTIPLE_MYELOMA_CD1_UP



features = 15 , max = 2