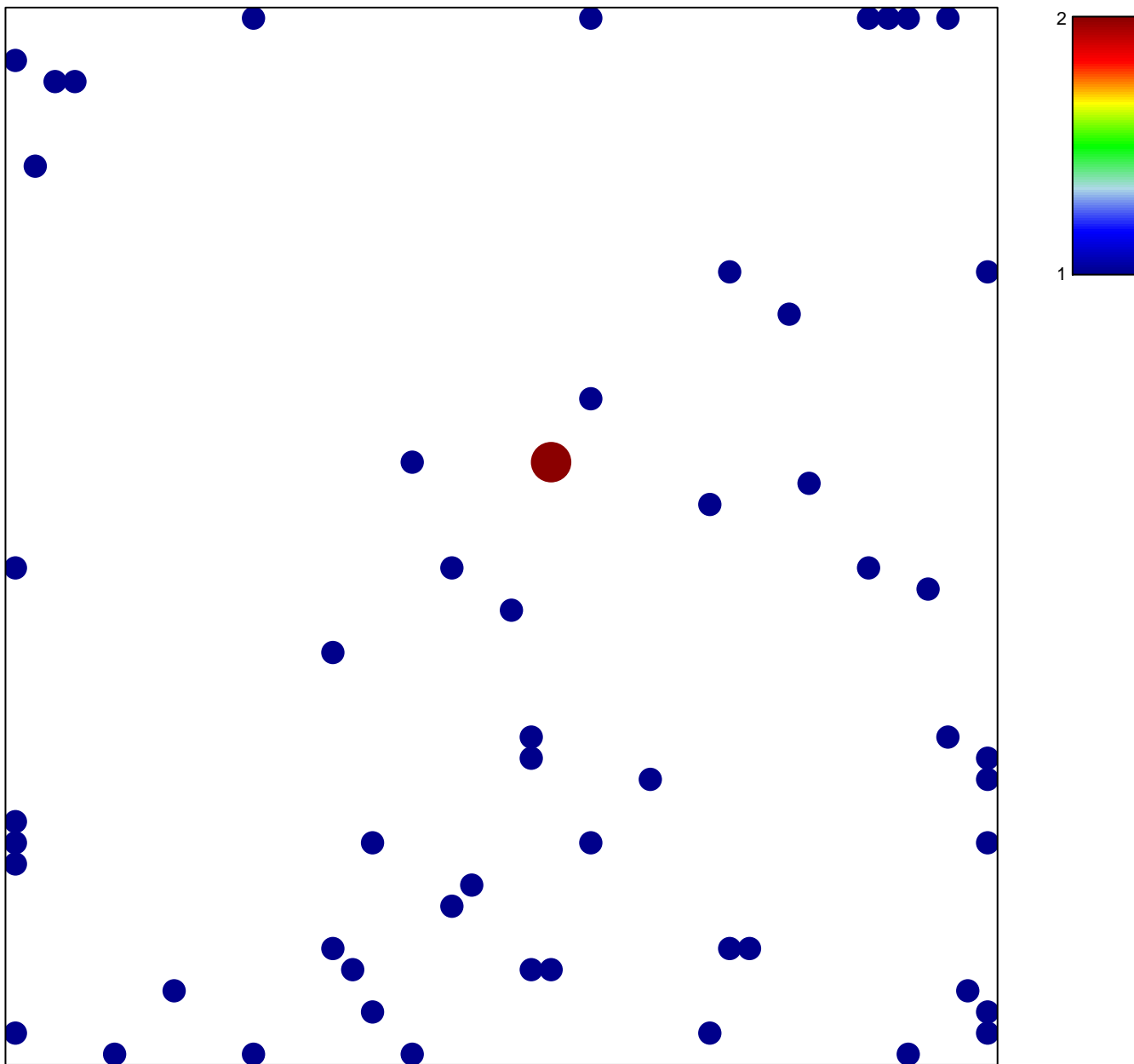
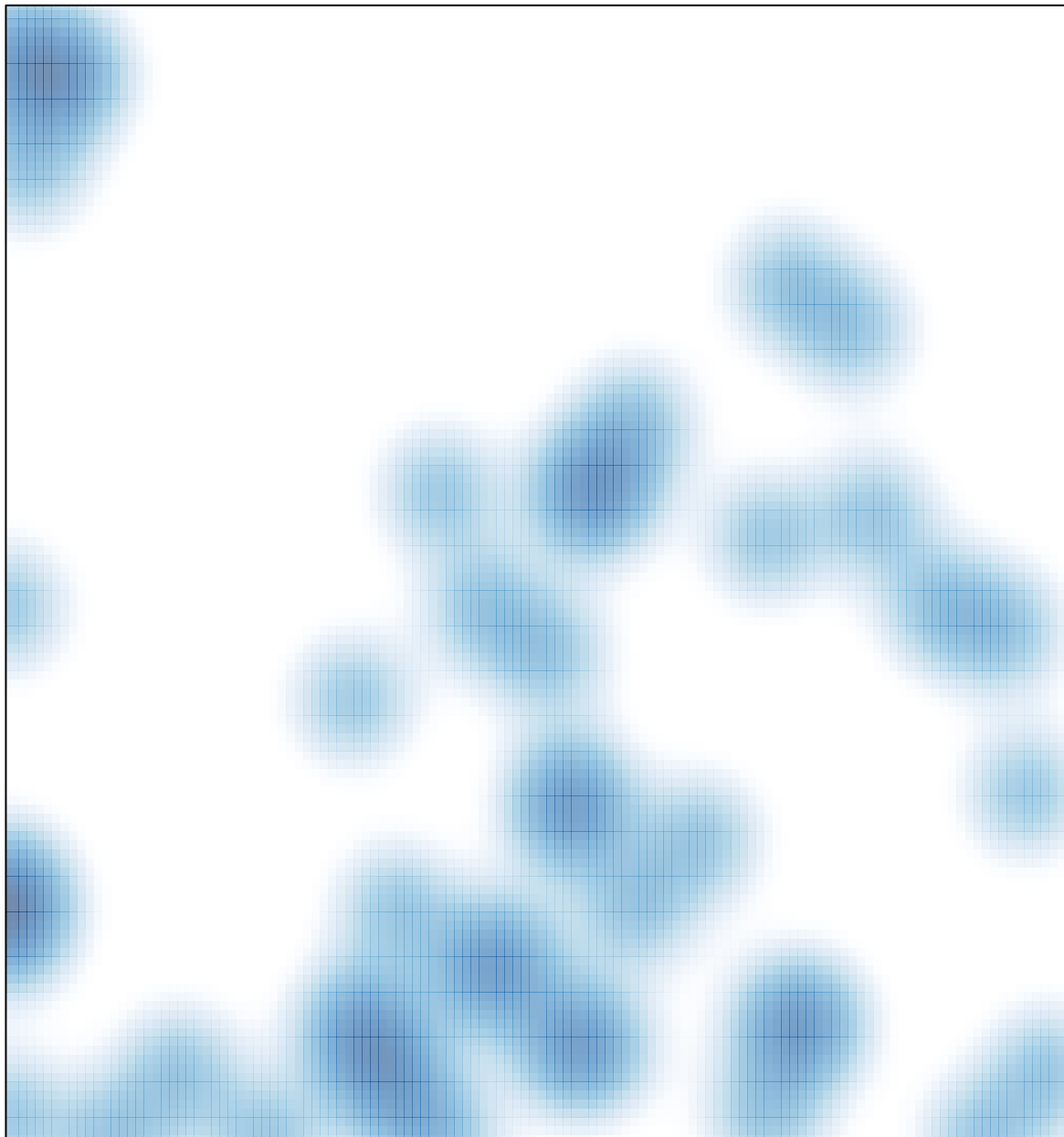


ZHAN_MULTIPLE_MYELOMA_CD1_VS_CD2_UP



features = 56
chi-square p = 0.8

ZHAN_MULTIPLE_MYELOMA_CD1_VS_CD2_UP



features = 56 , max = 2