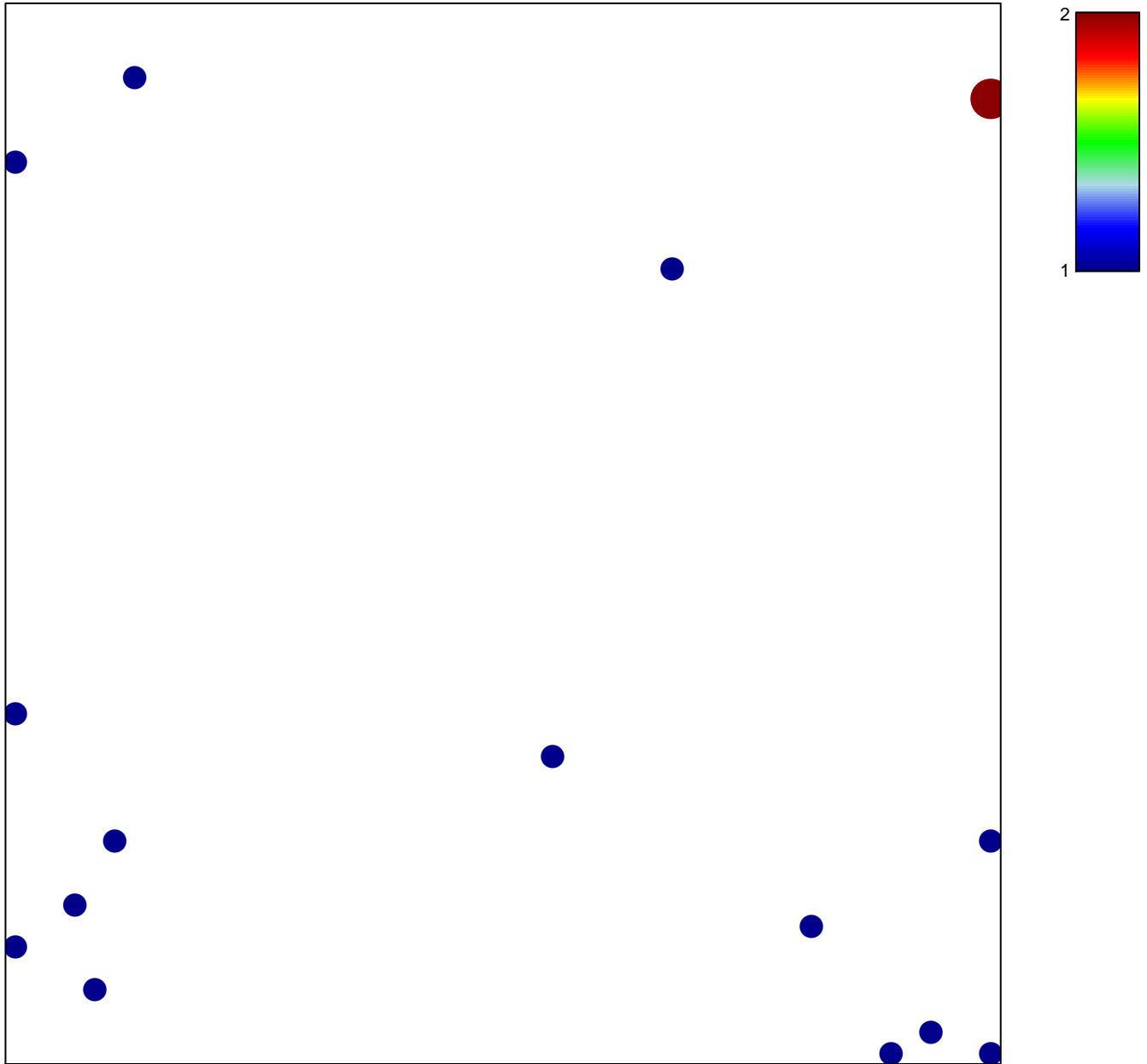
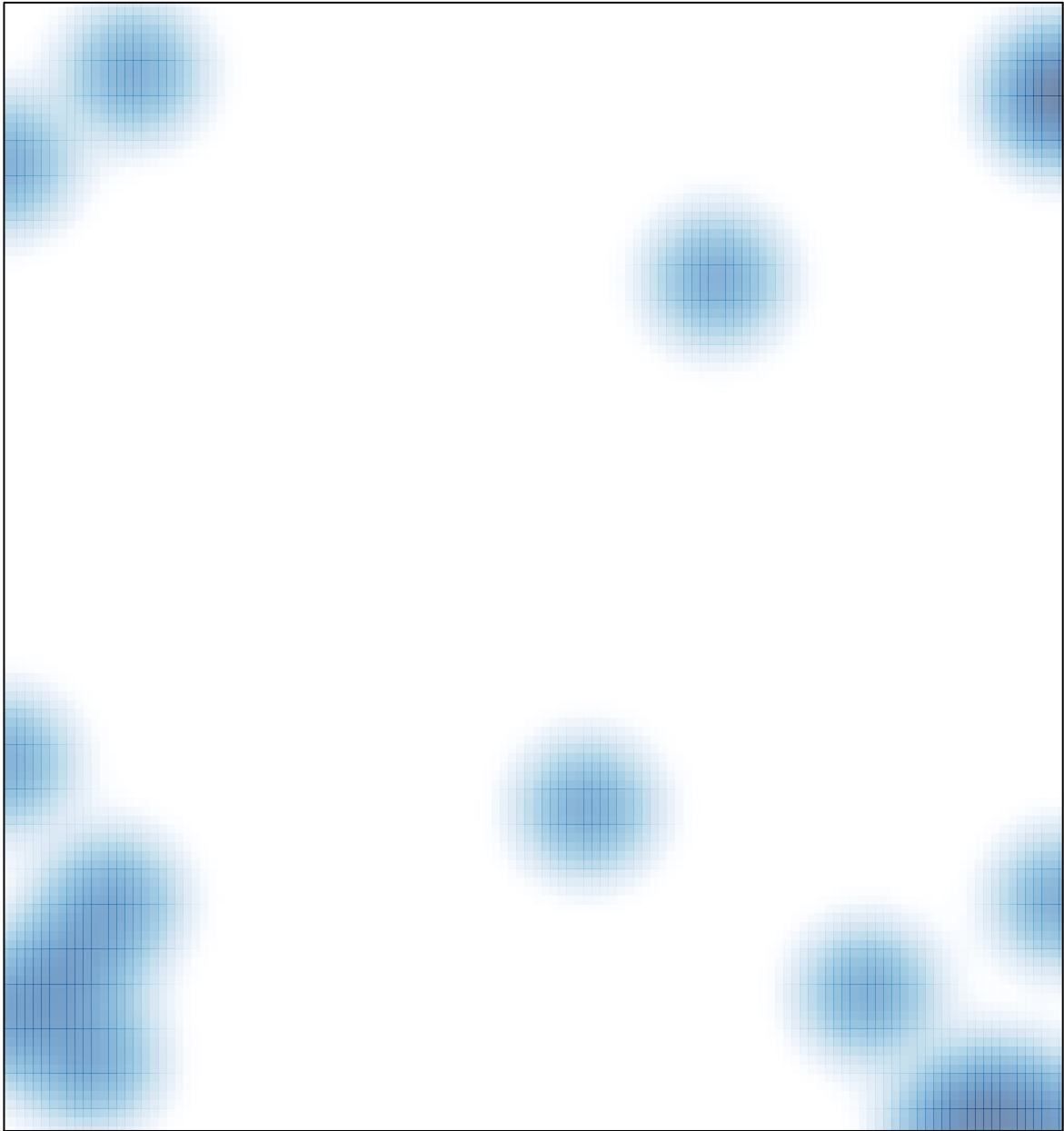


ZHAN_MULTIPLE_MYELOMA_CD1_VS_CD2_UP



features = 16
chi-square p = 0.81

ZHAN_MULTIPLE_MYELOMA_CD1_VS_CD2_UP



features = 16 , max = 2