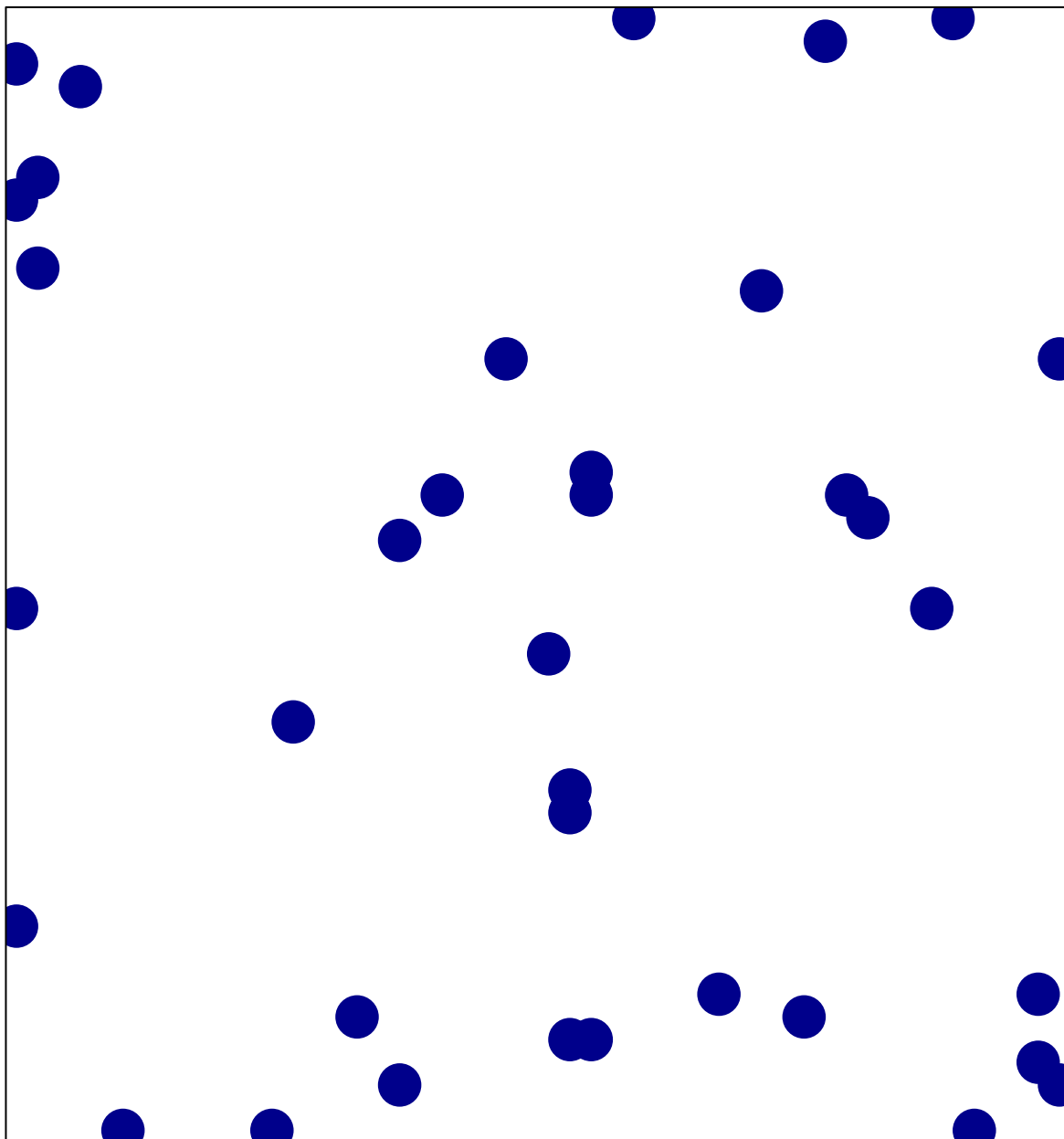
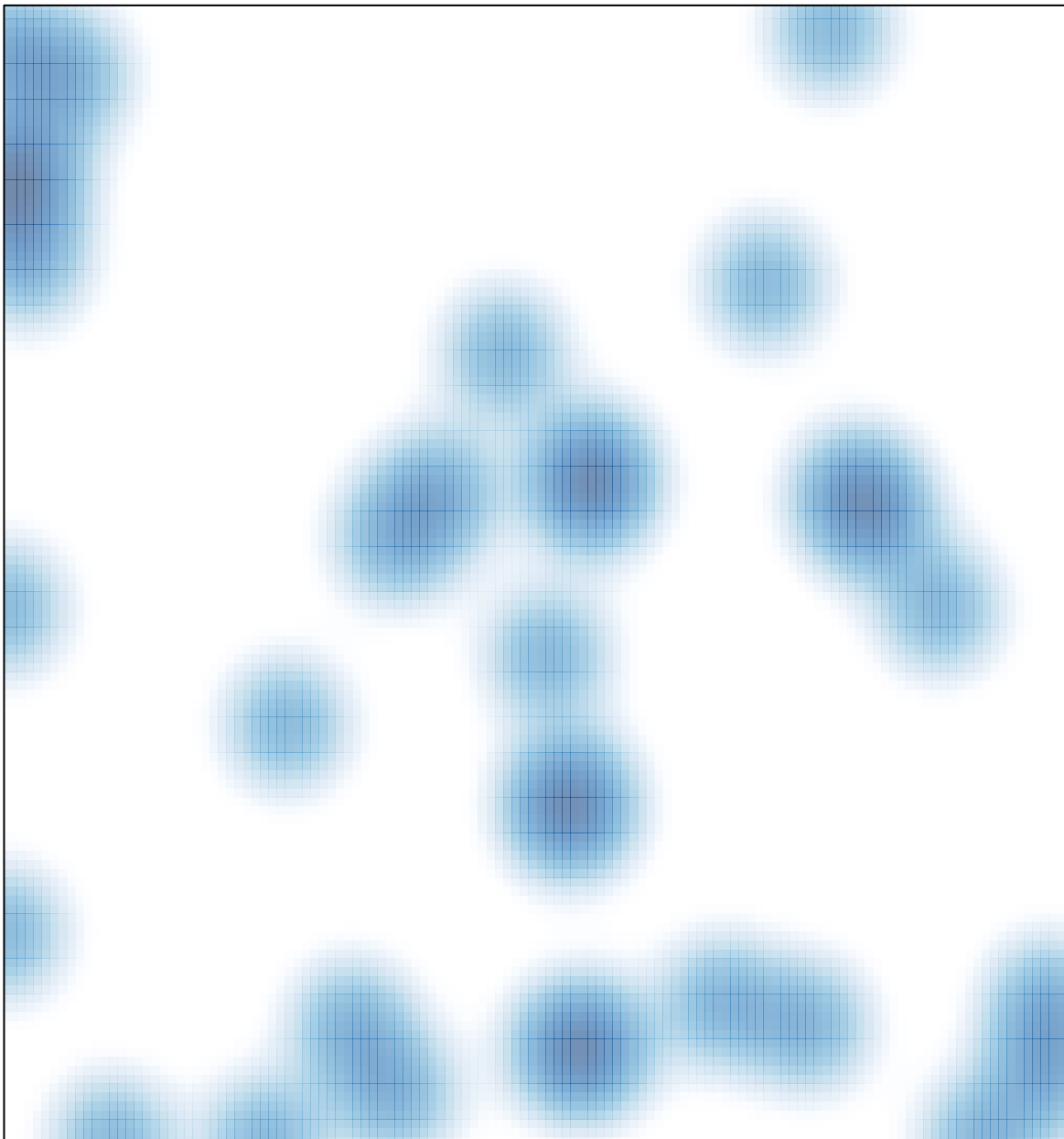


ZHAN_MULTIPLE_MYELOMA_CD1_UP



features = 36
chi-square p = 0.83

ZHAN_MULTIPLE_MYELOMA_CD1_UP



features = 36 , max = 1