CORRE_MULTIPLE_MYELOMA_UP
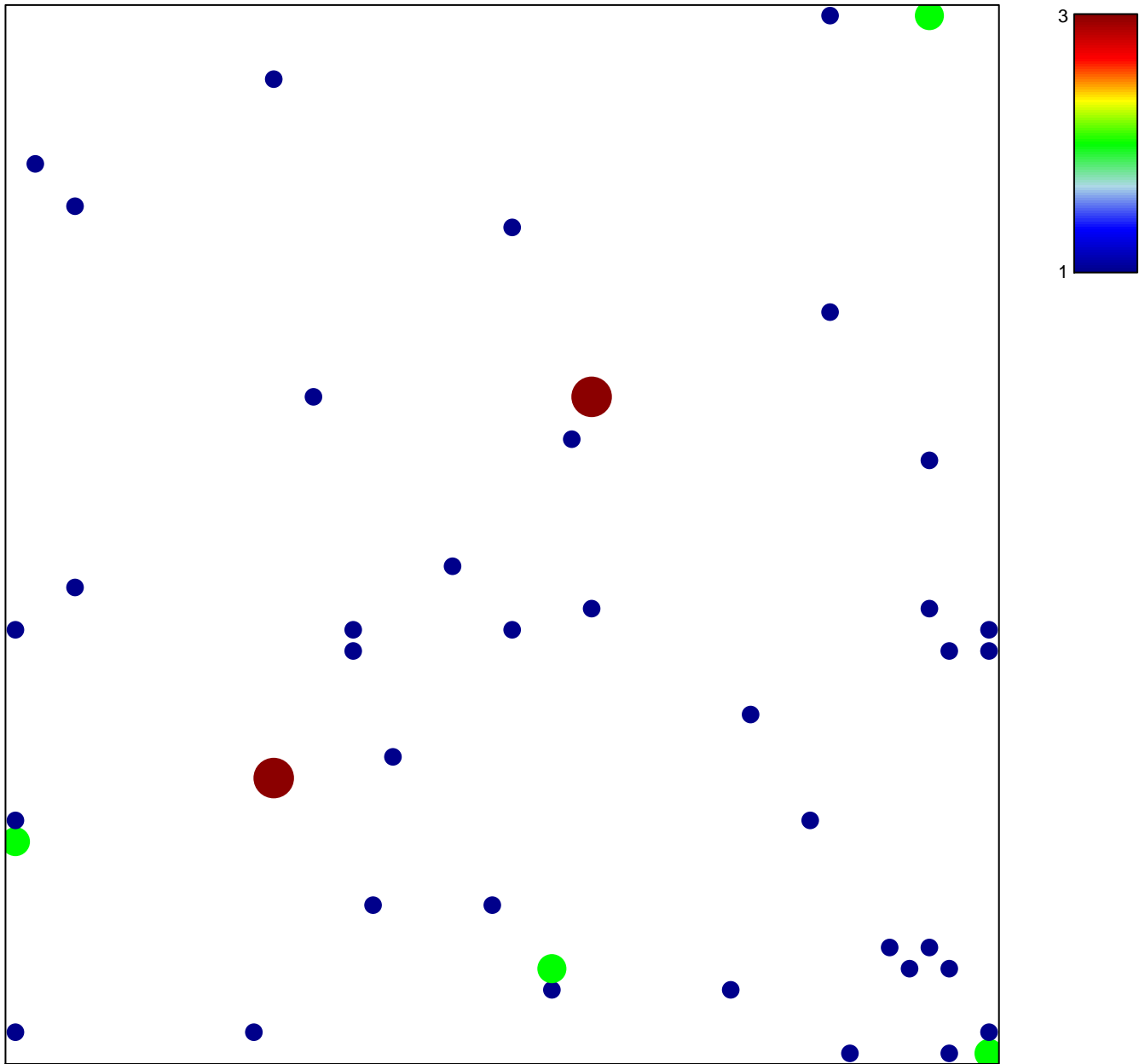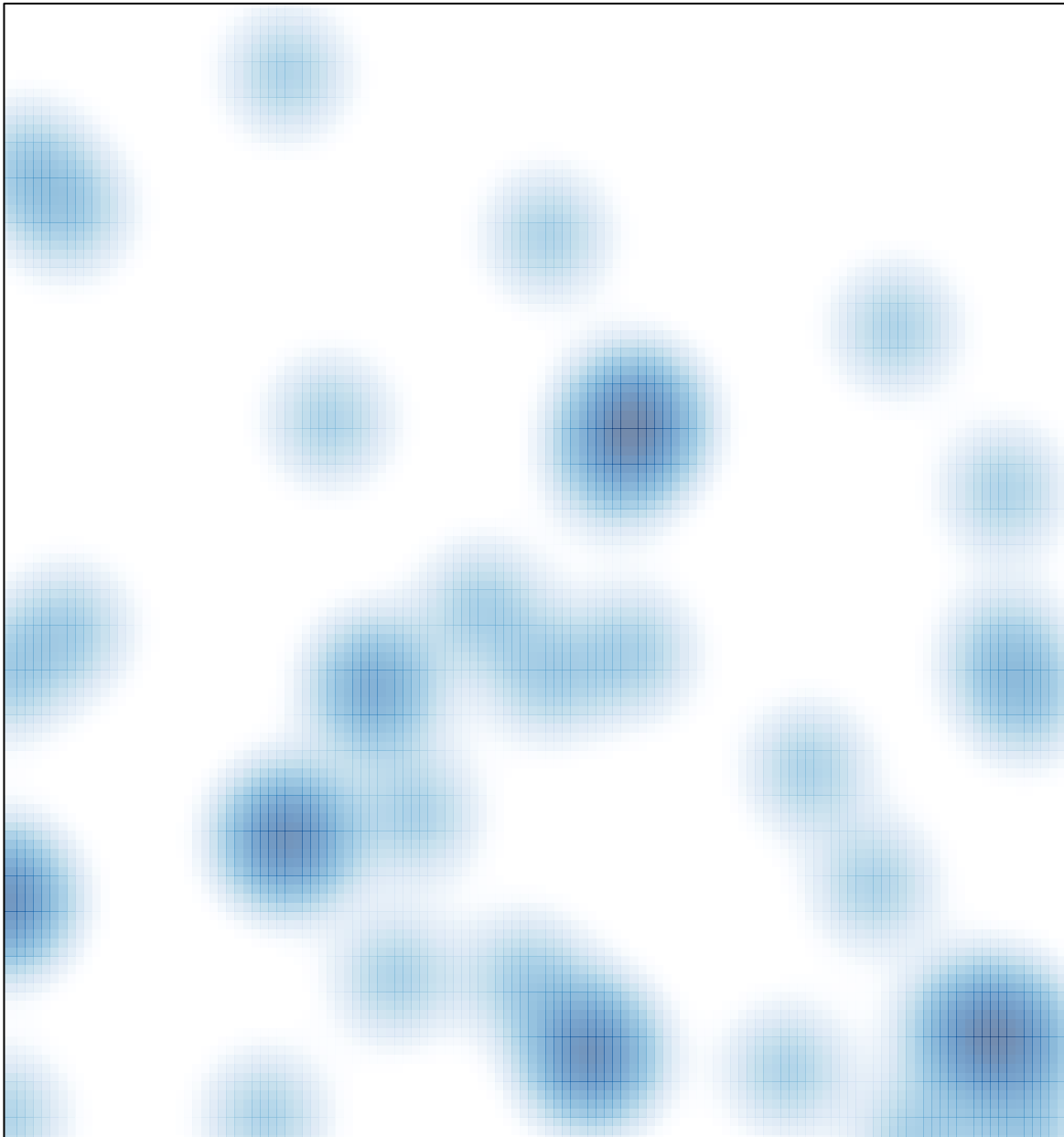

features = 51

chi−square p = 0.79

CORRE_MULTIPLE_MYELOMA_UP

features = 51 , max = 3