CORRE_MULTIPLE_MYELOMA_DN
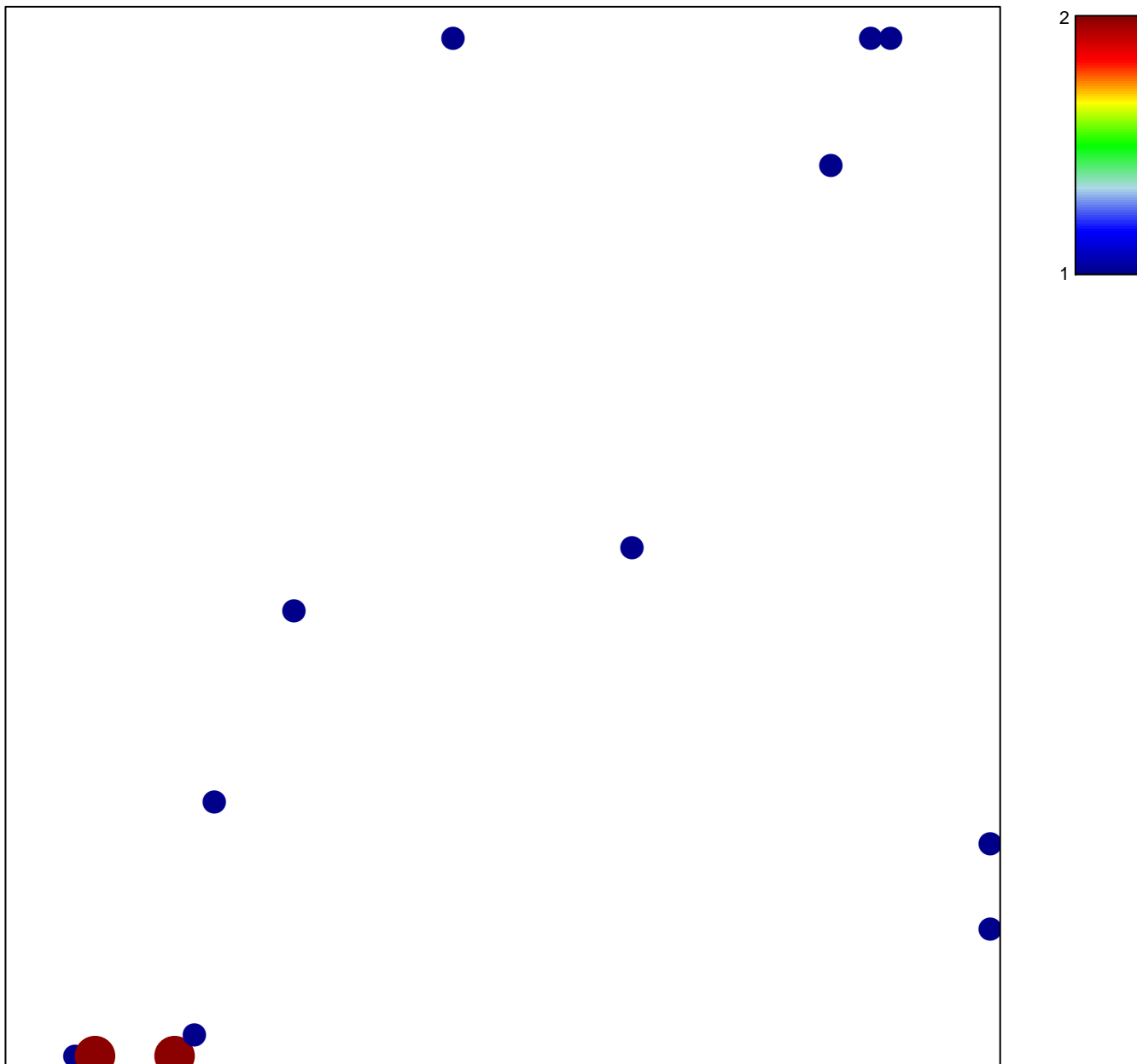

features = 15
chi−square p = 0.79

CORRE_MULTIPLE_MYELOMA_DN

features = 15 , max = 2