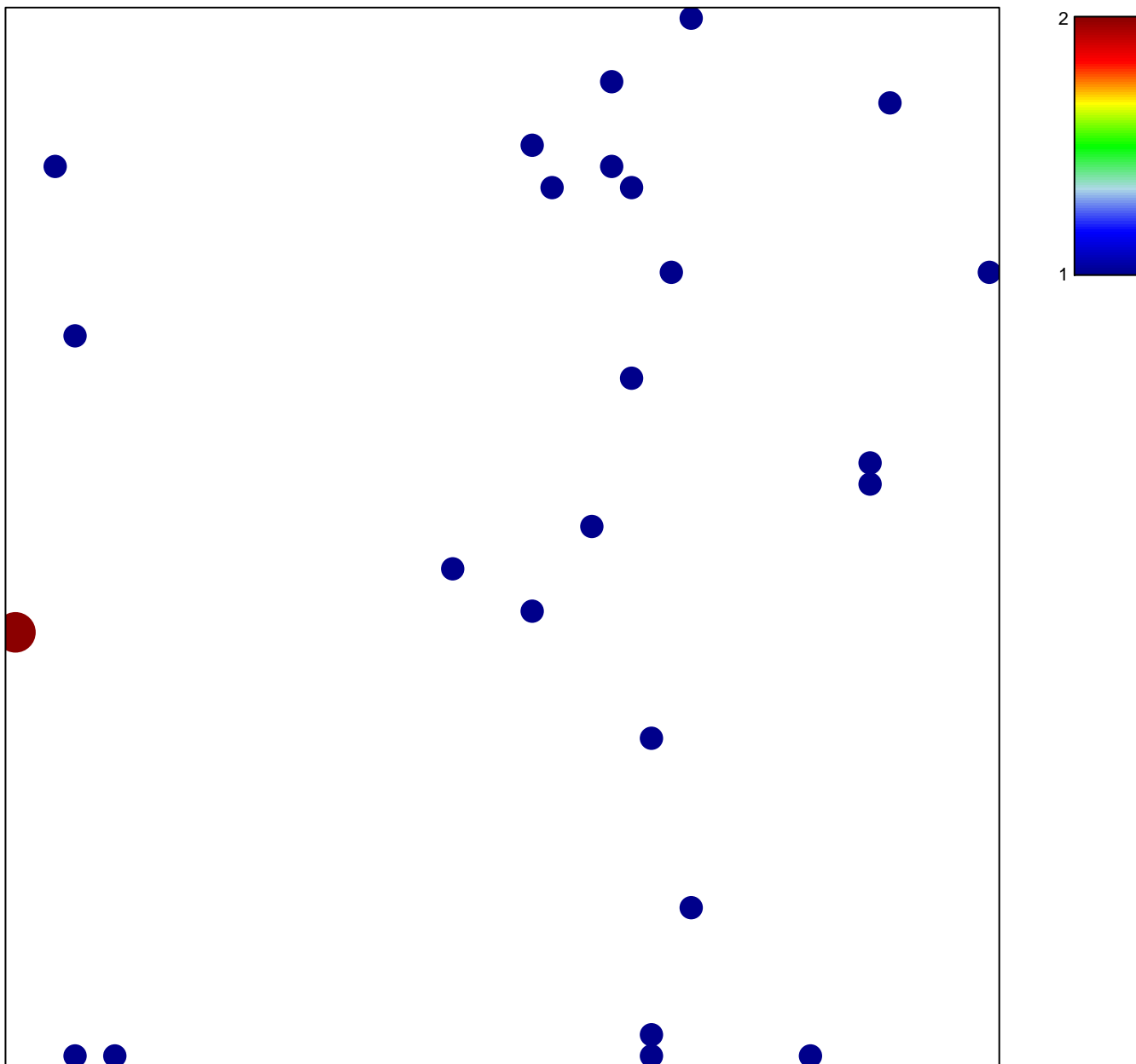


# BILANGES\_SERUM\_AND\_RAPAMYCIN\_SENSITIVE\_GENES



# features = 26  
chi-square p = 0.82

# ANGES\_SERUM\_AND\_RAPAMYCIN\_SENSITIVE\_GE



# features = 26 , max = 2