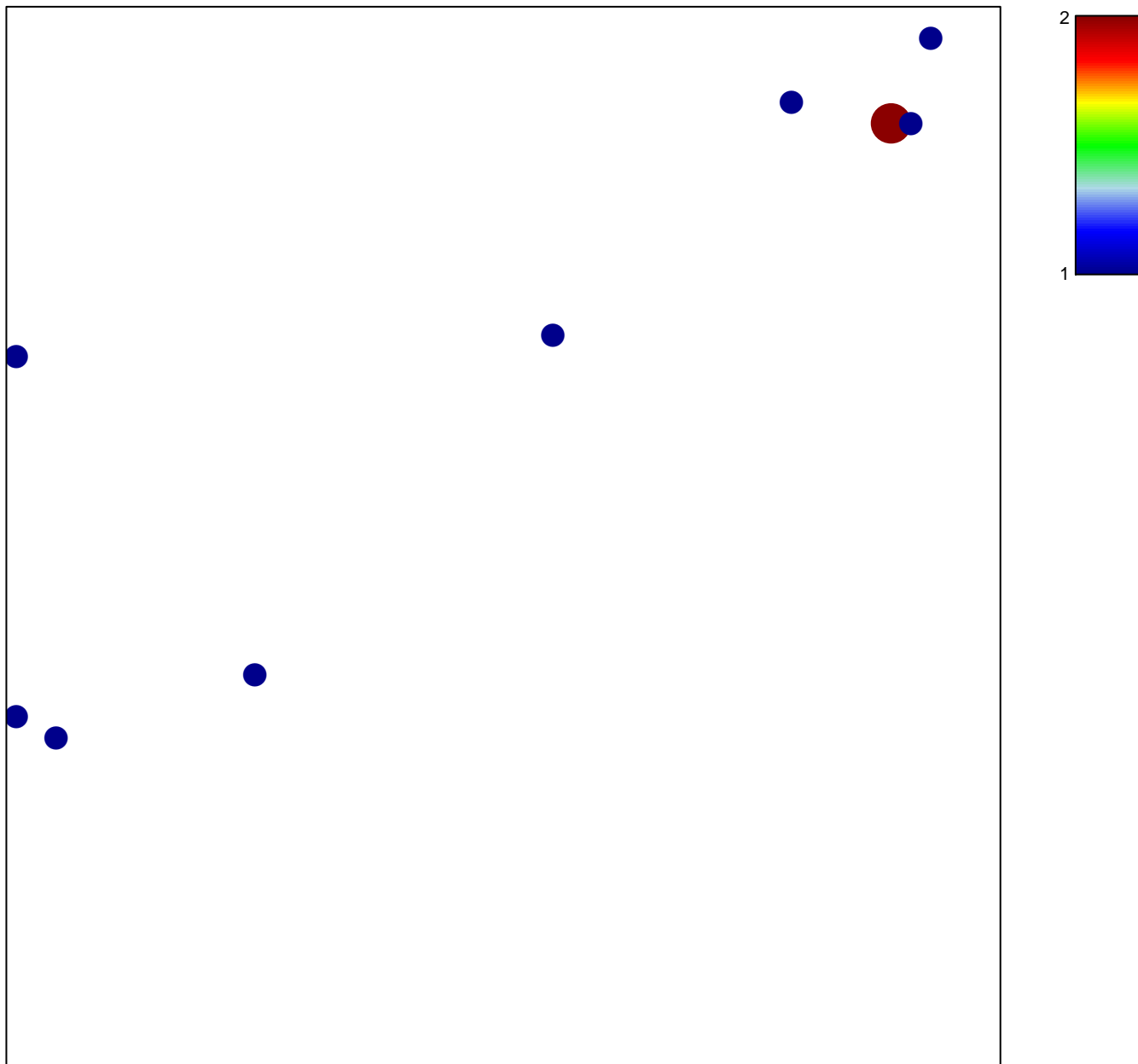


# REACTOME\_GLOBAL\_GENOMIC\_NER



# features = 10  
chi-square p = 0.82

# REACTOME\_GLOBAL\_GENOMIC\_NER



# features = 10 , max = 2