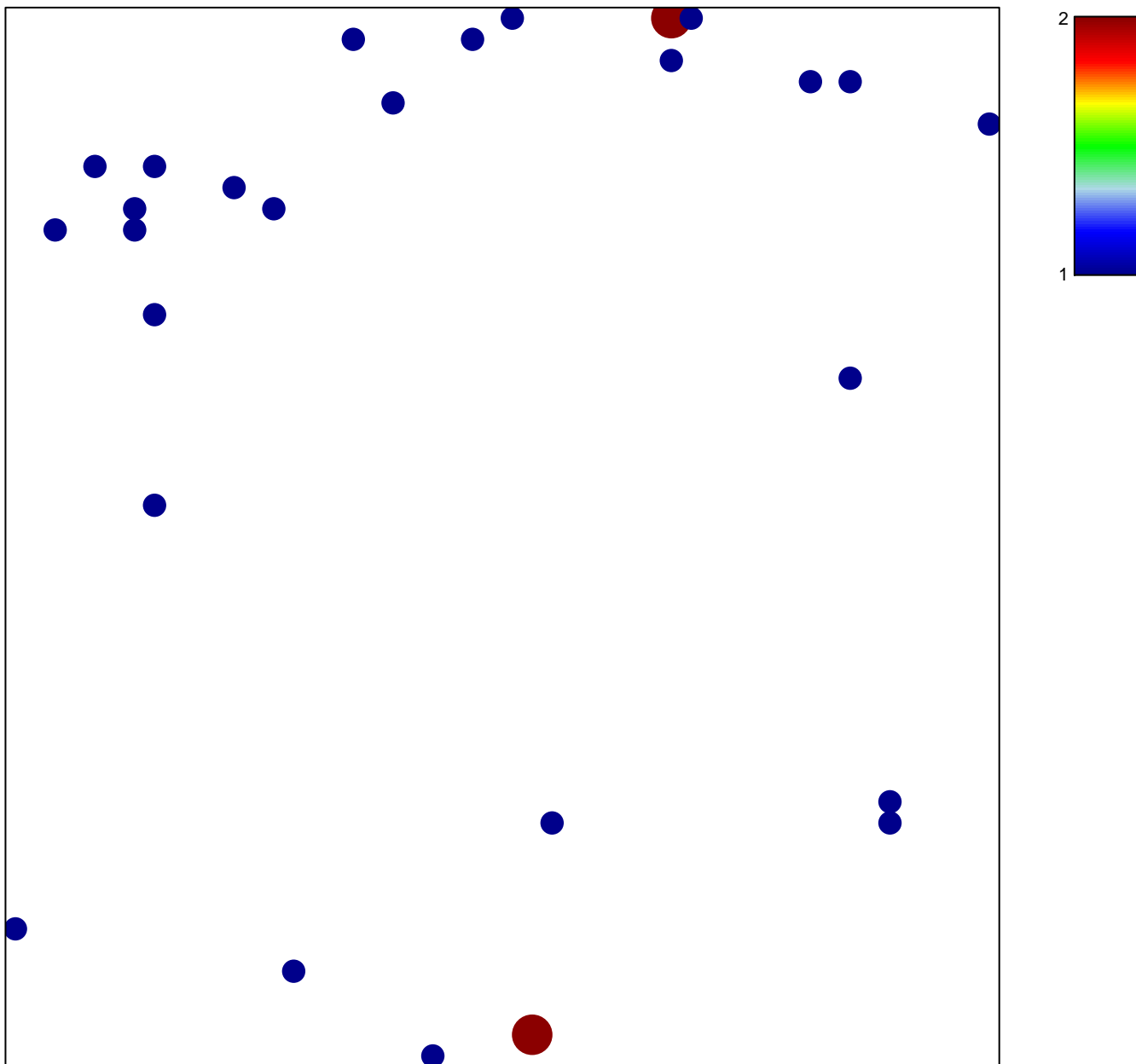


# spliceosomal snRNP assembly



# features = 29  
chi-square p = 0.83

# spliceosomal snRNP assembly



# features = 29 , max = 2