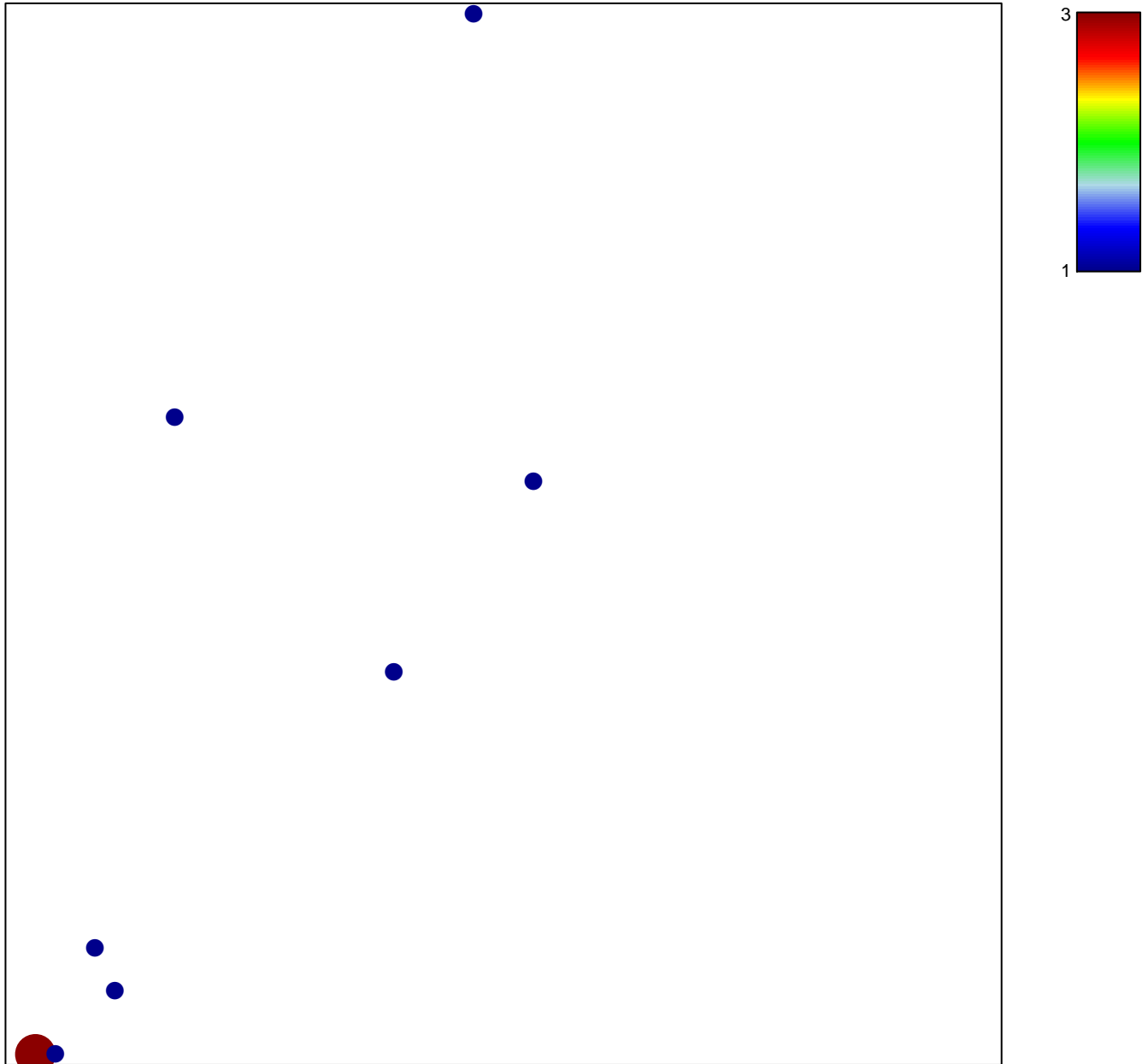


# protein heterotrimerization



# features = 10  
chi-square p = 0.79

# protein heterotrimerization



# features = 10 , max = 3