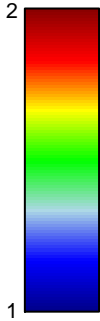
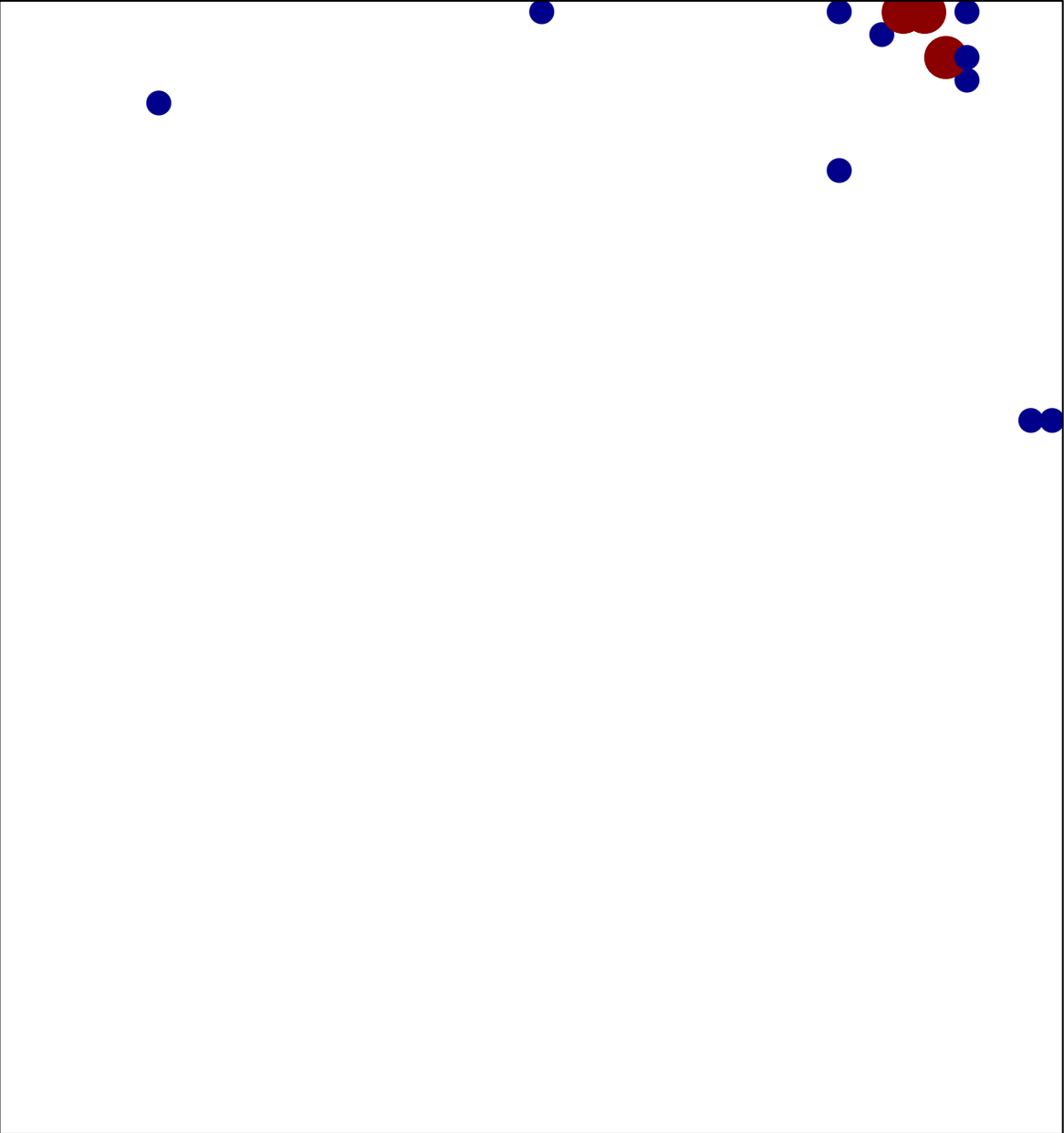


# WOLFER\_overlap genes



# features = 16  
chi-square p = 0.74

# WOLFER\_overlap genes



# features = 16 , max = 2