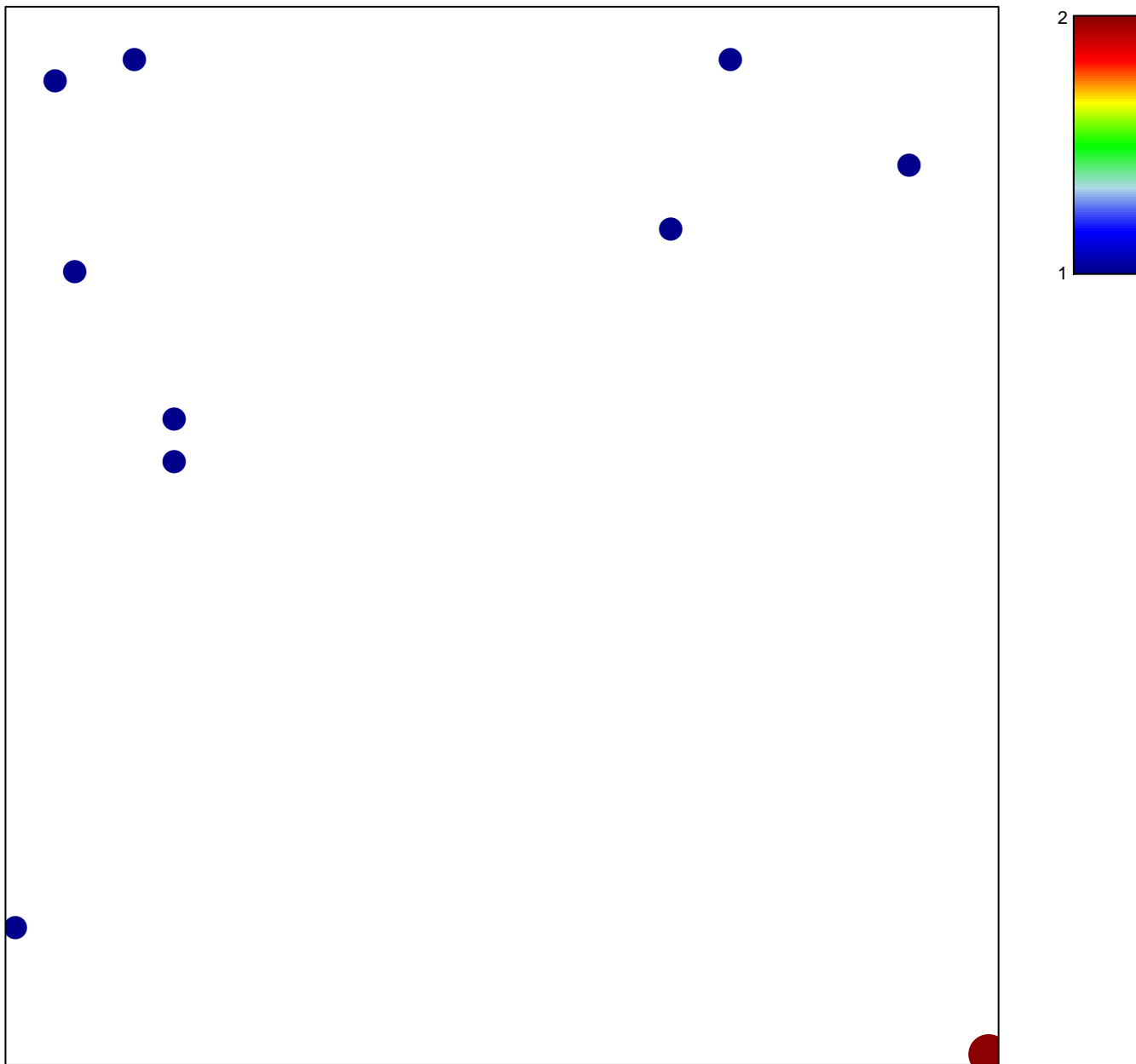
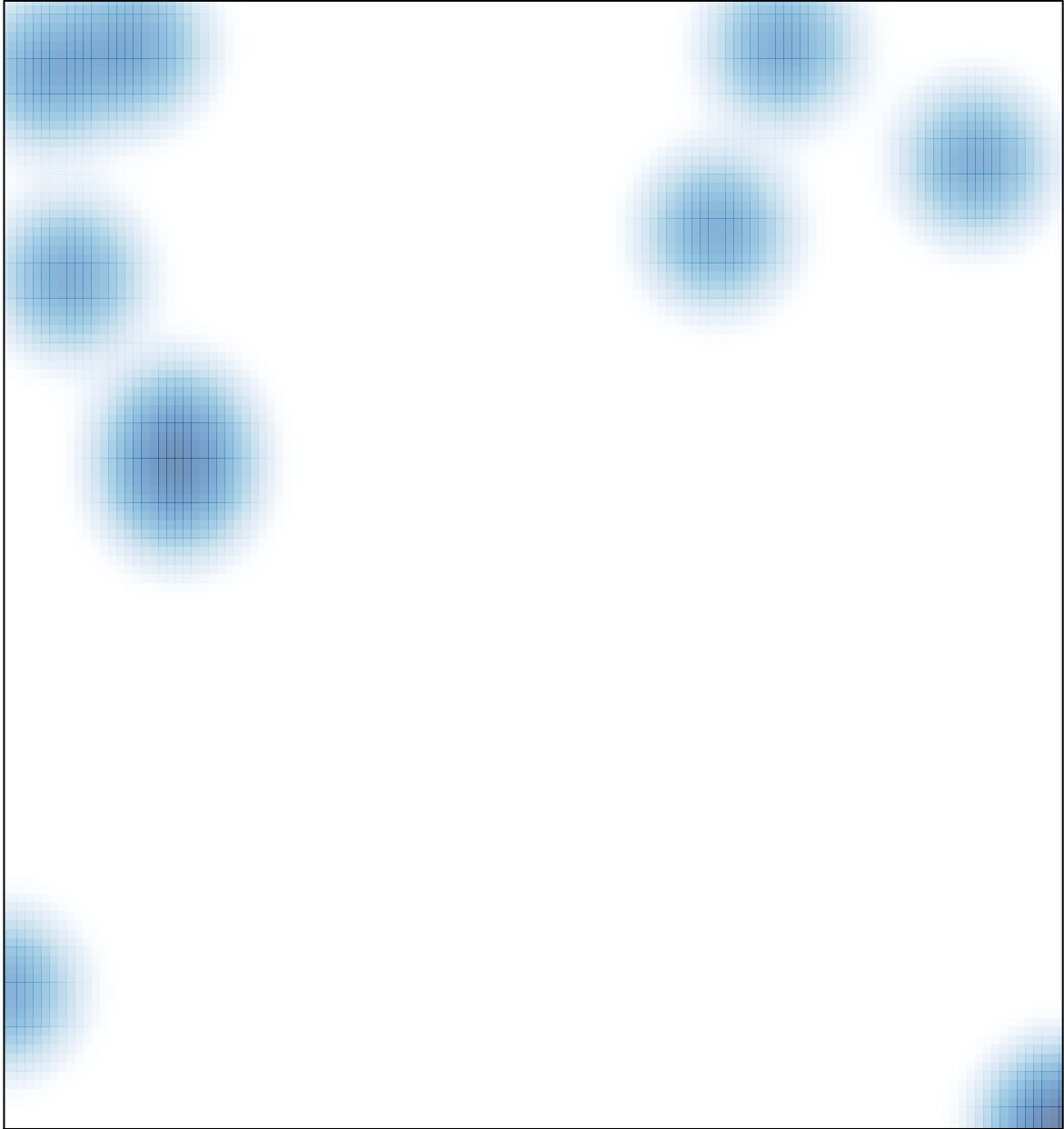


# chaperone mediated protein folding requiring cofactor



# features = 11  
chi-square p = 0.83

# chaperone mediated protein folding requiring cofactors



# features = 11 , max = 2