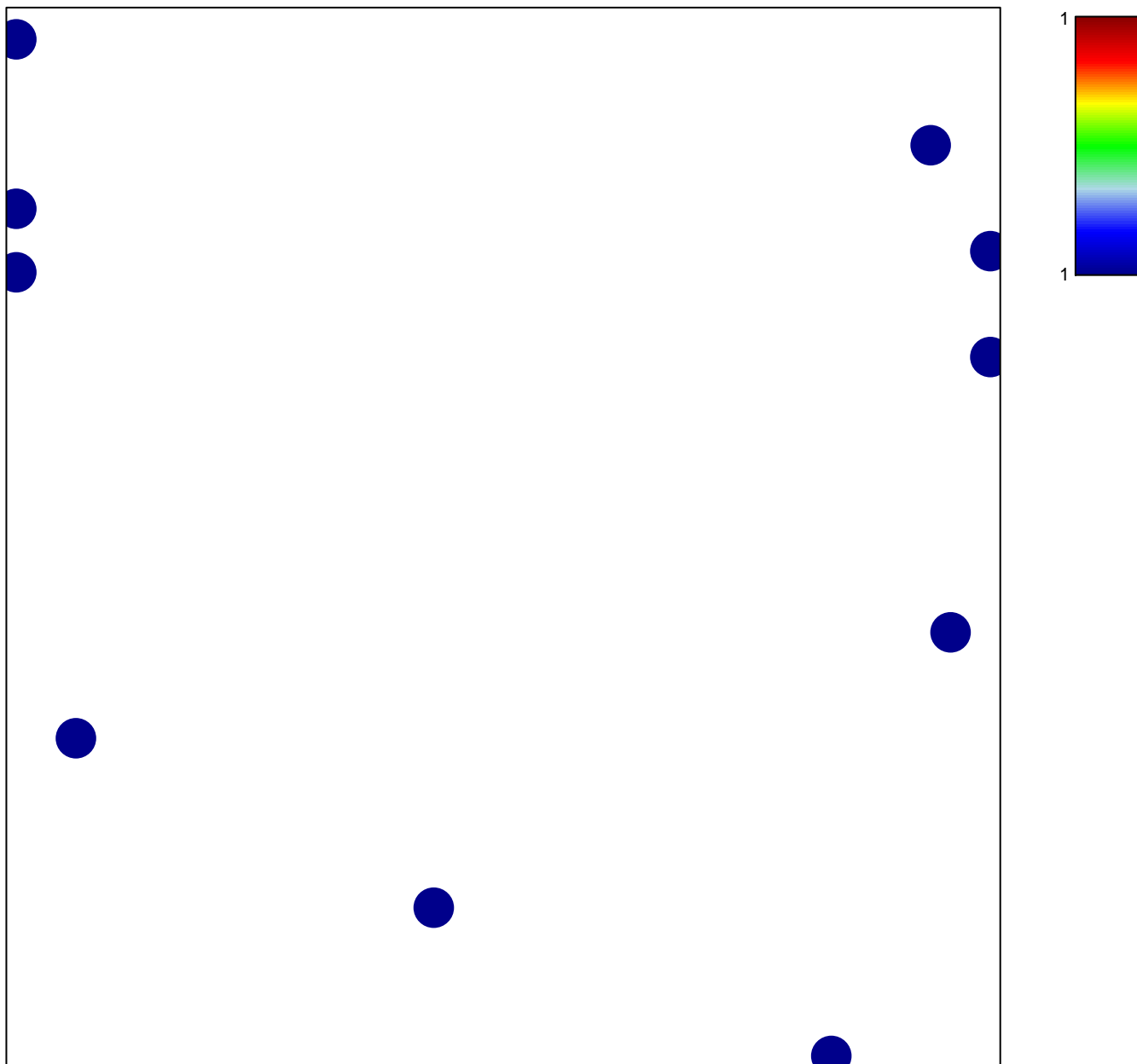
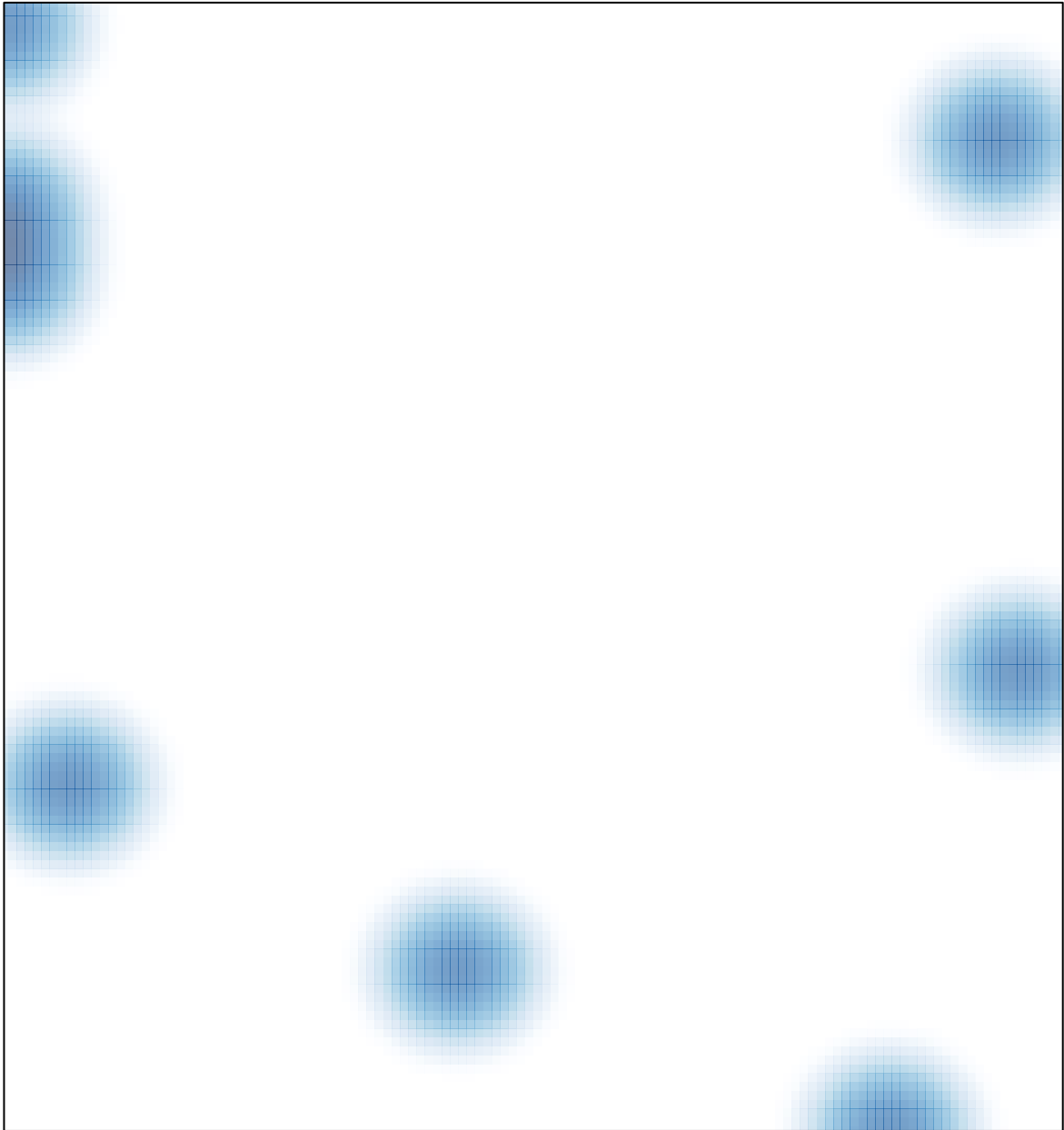


# protein deacetylation



# features = 10  
chi-square p = 0.83

# protein deacetylation



# features = 10 , max = 1