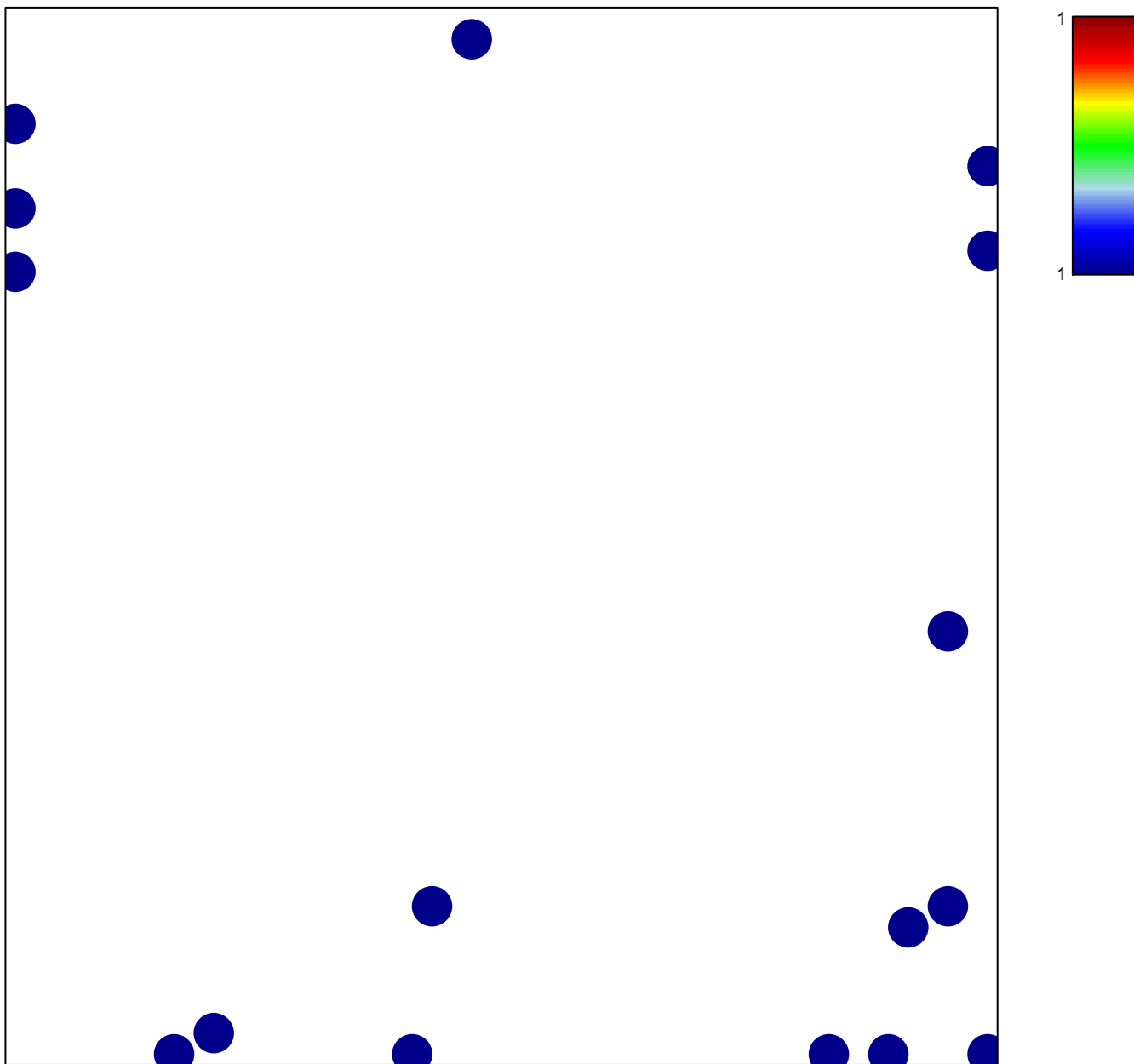
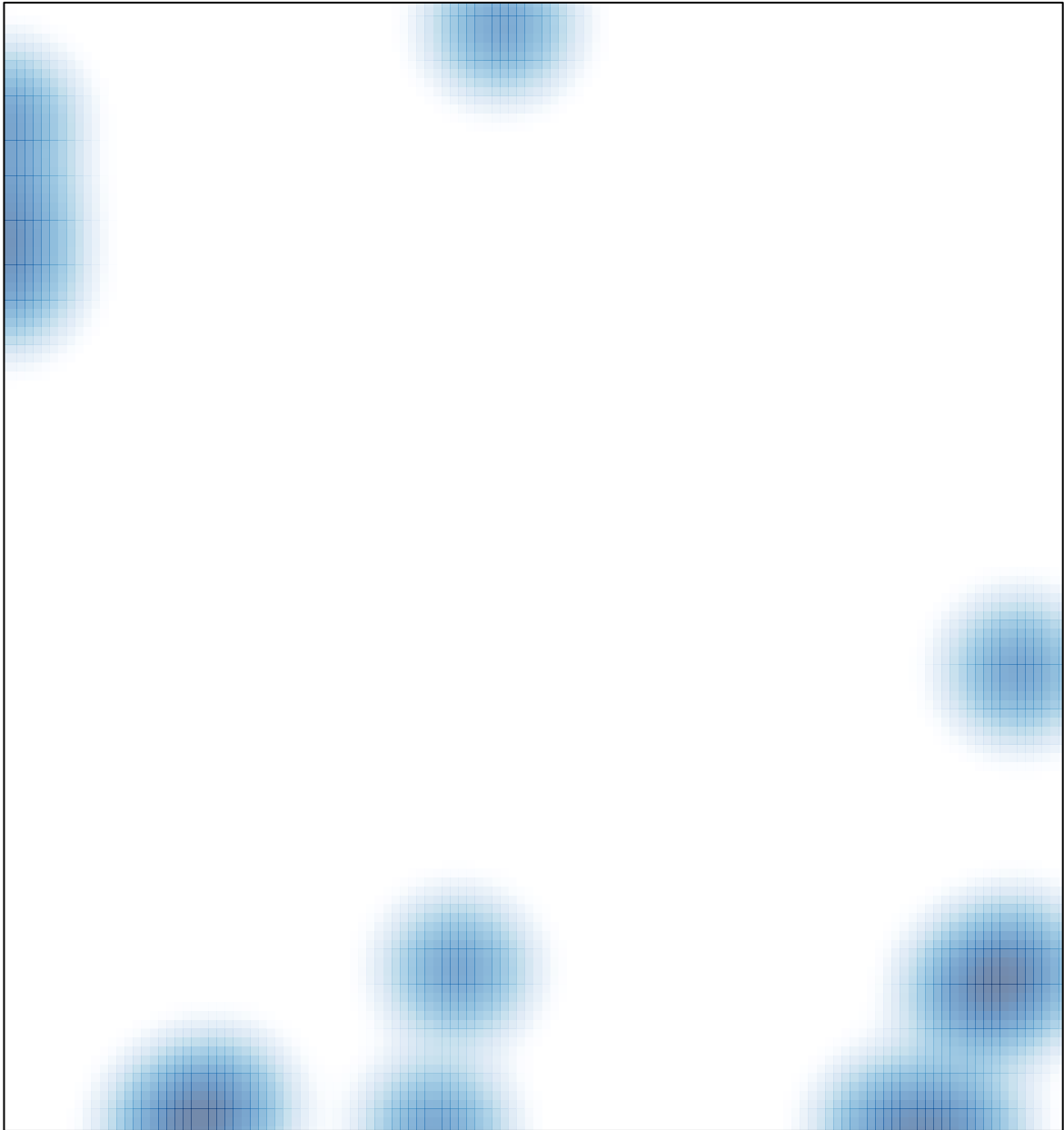


# histone H4 deacetylation



# features = 16  
chi-square p = 0.79

# histone H4 deacetylation



# features = 16 , max = 1