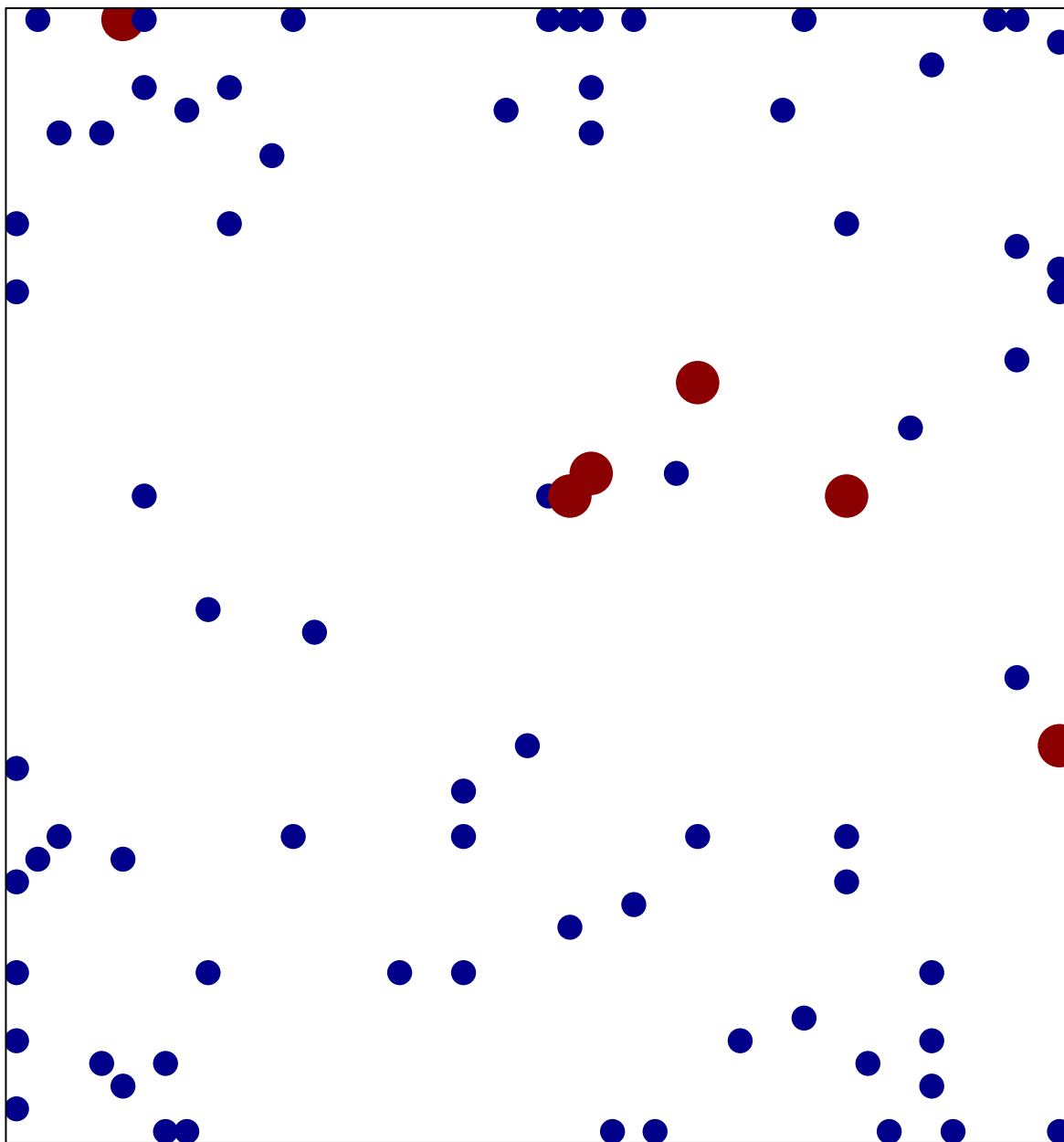
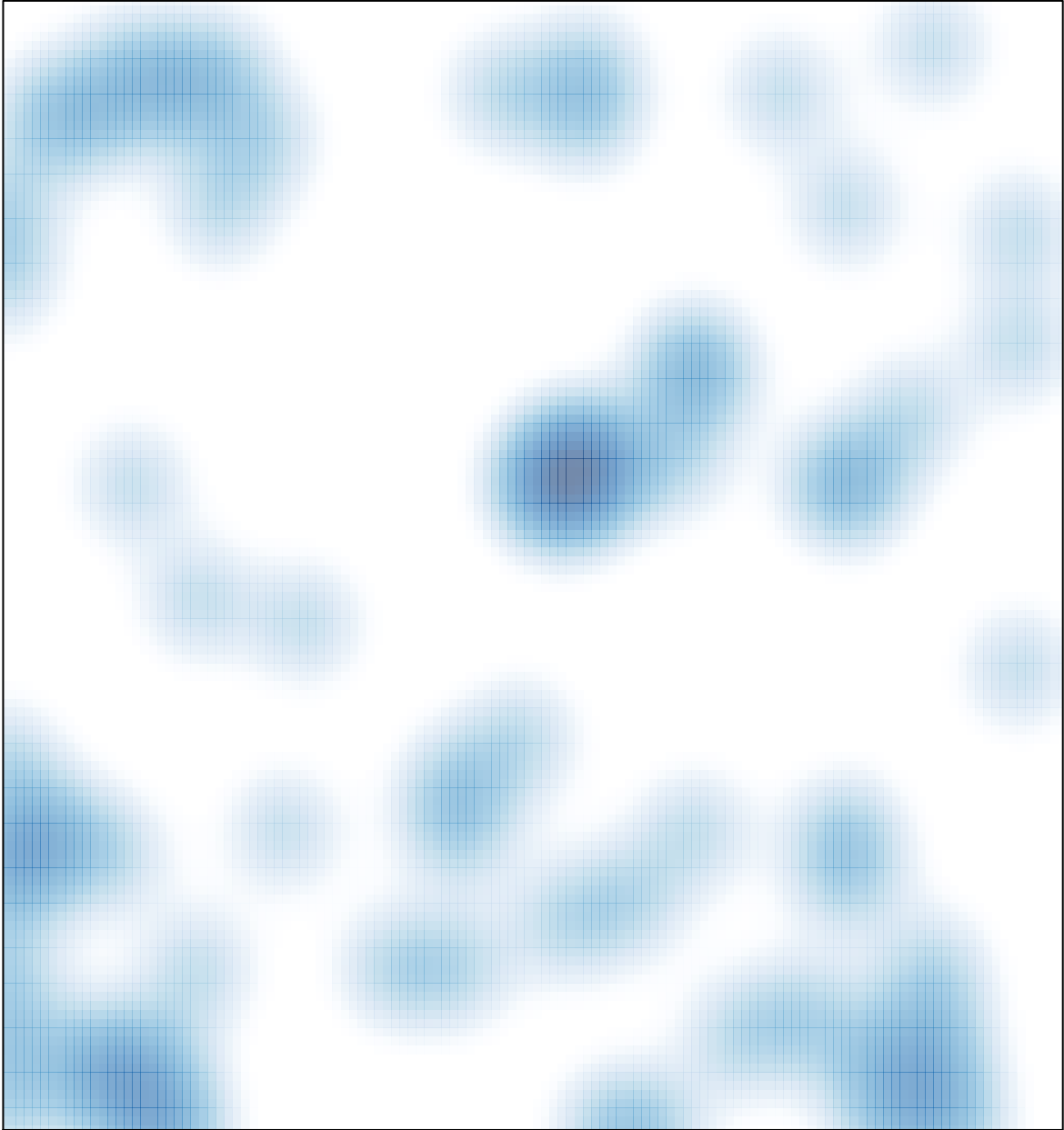


histone deacetylase binding



features = 85
chi-square p = 0.76

histone deacetylase binding



features = 85 , max = 2