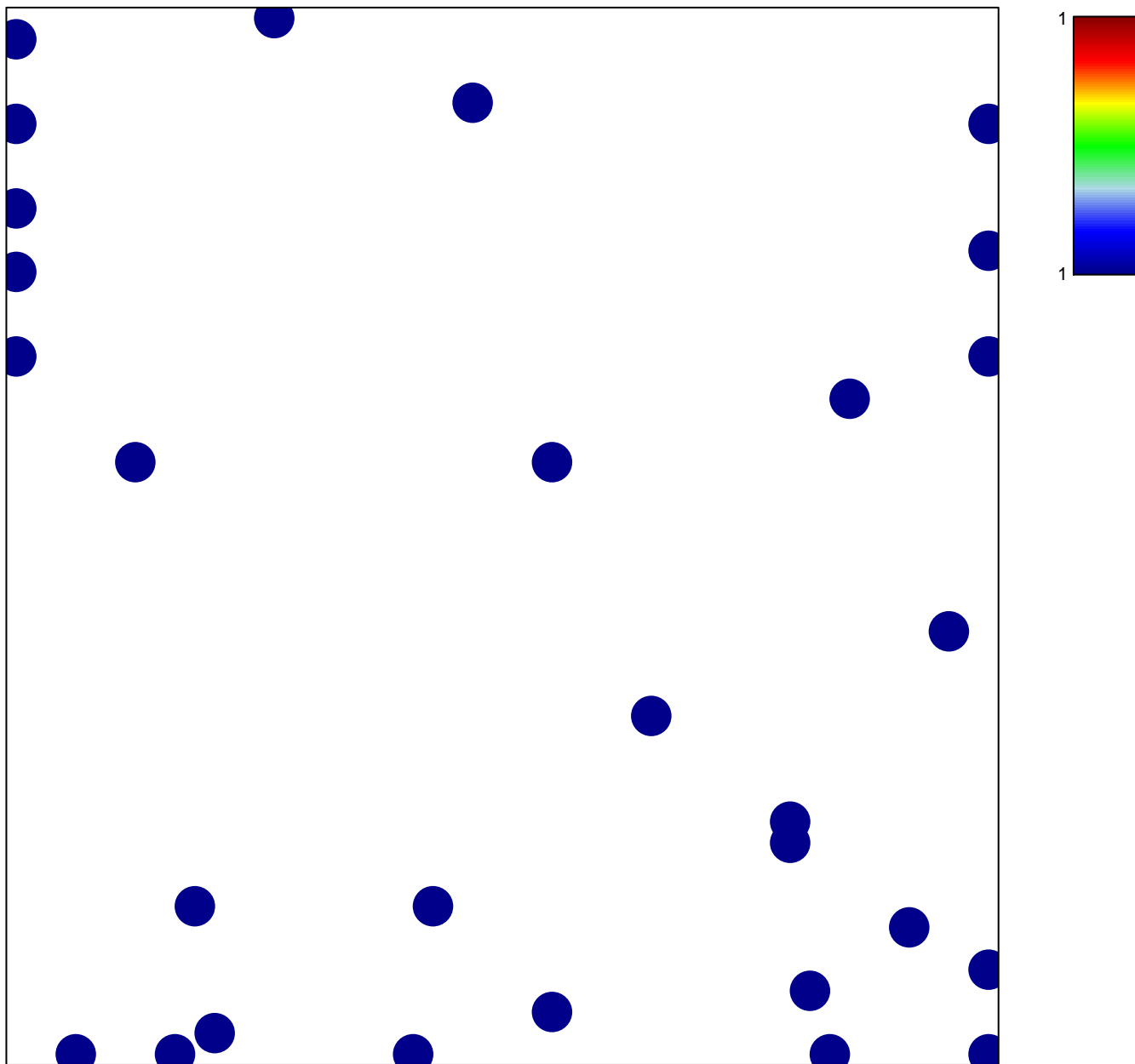
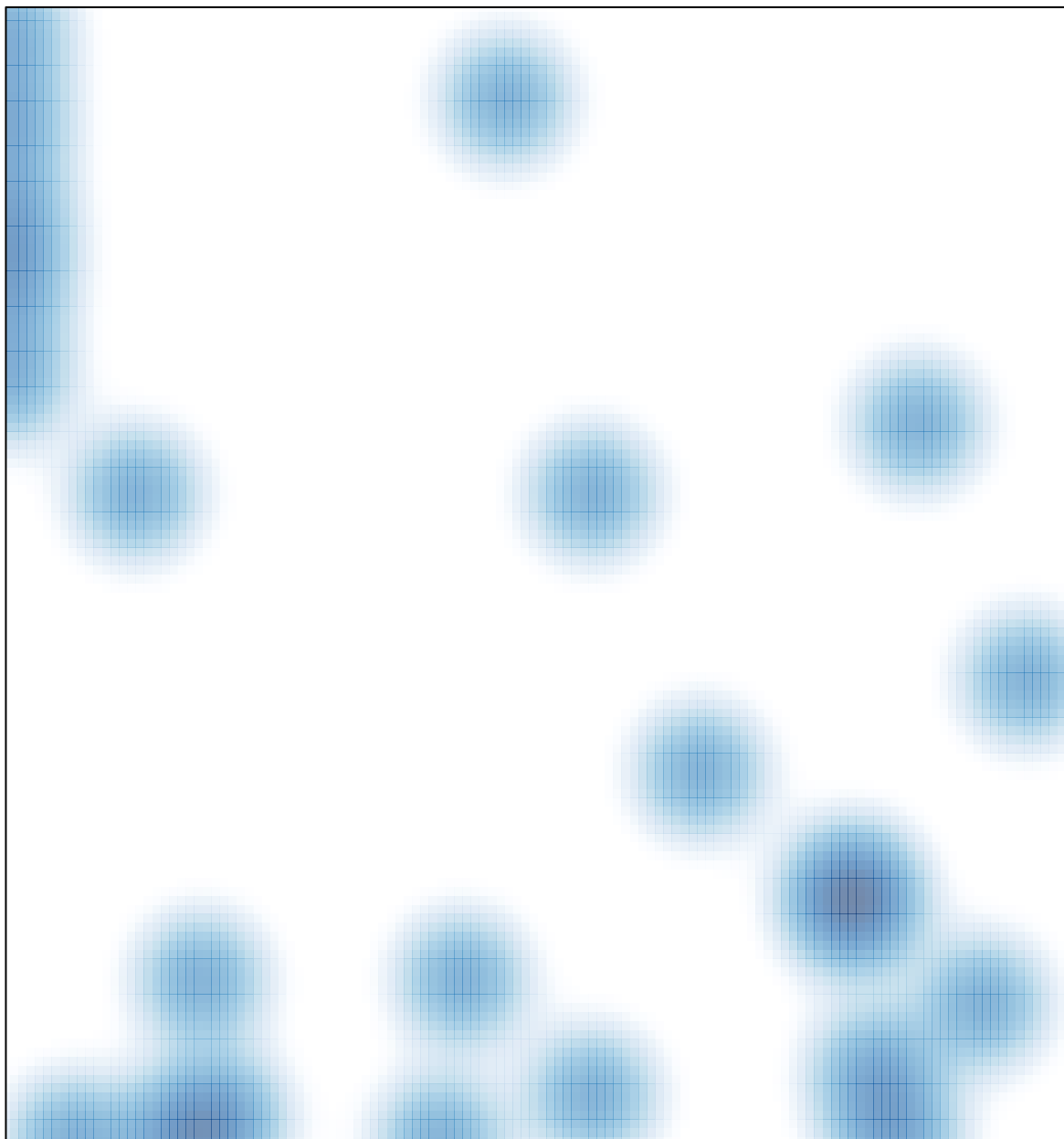


histone deacetylation



features = 29
chi-square p = 0.8

histone deacetylation



features = 29 , max = 1