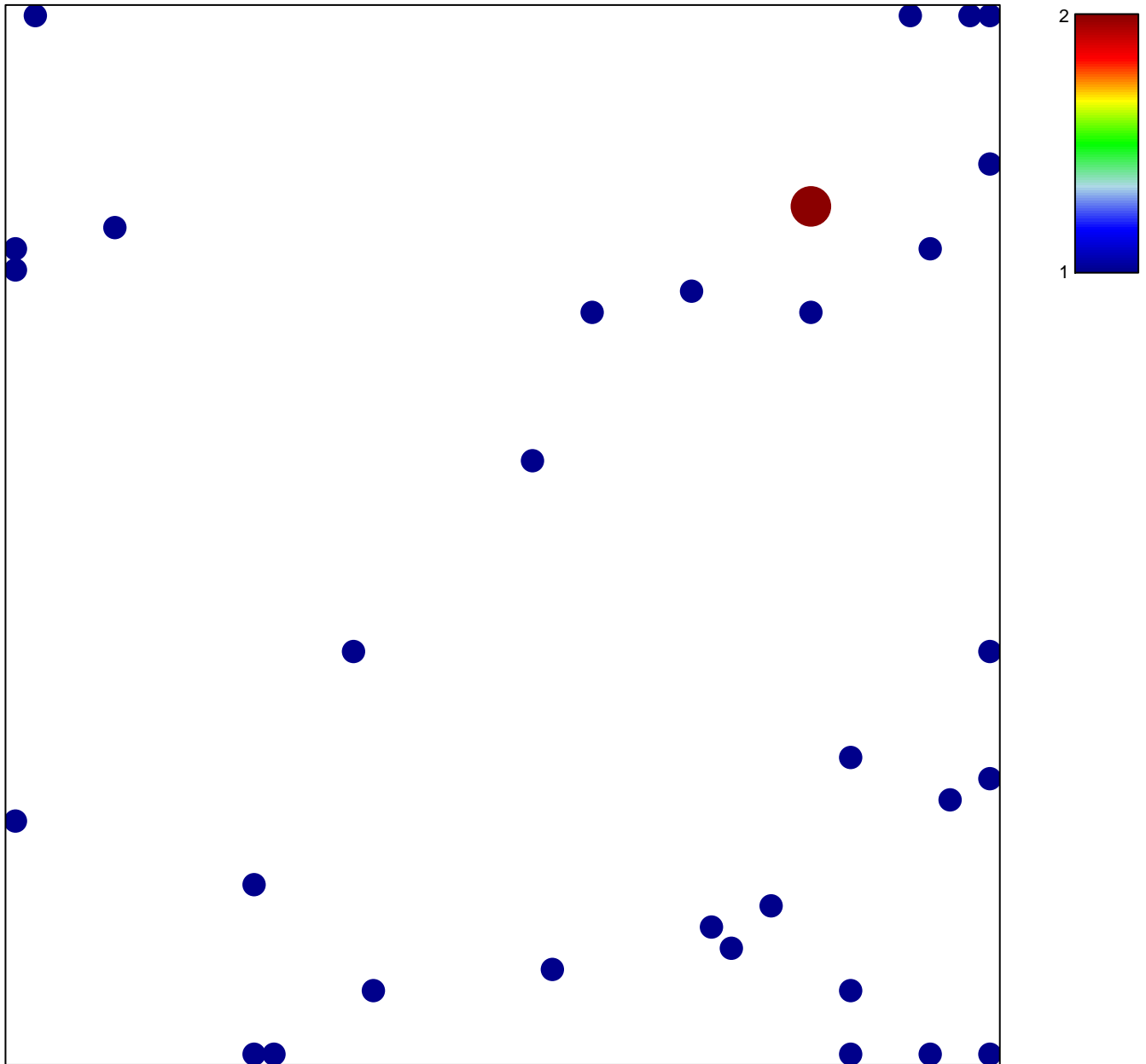
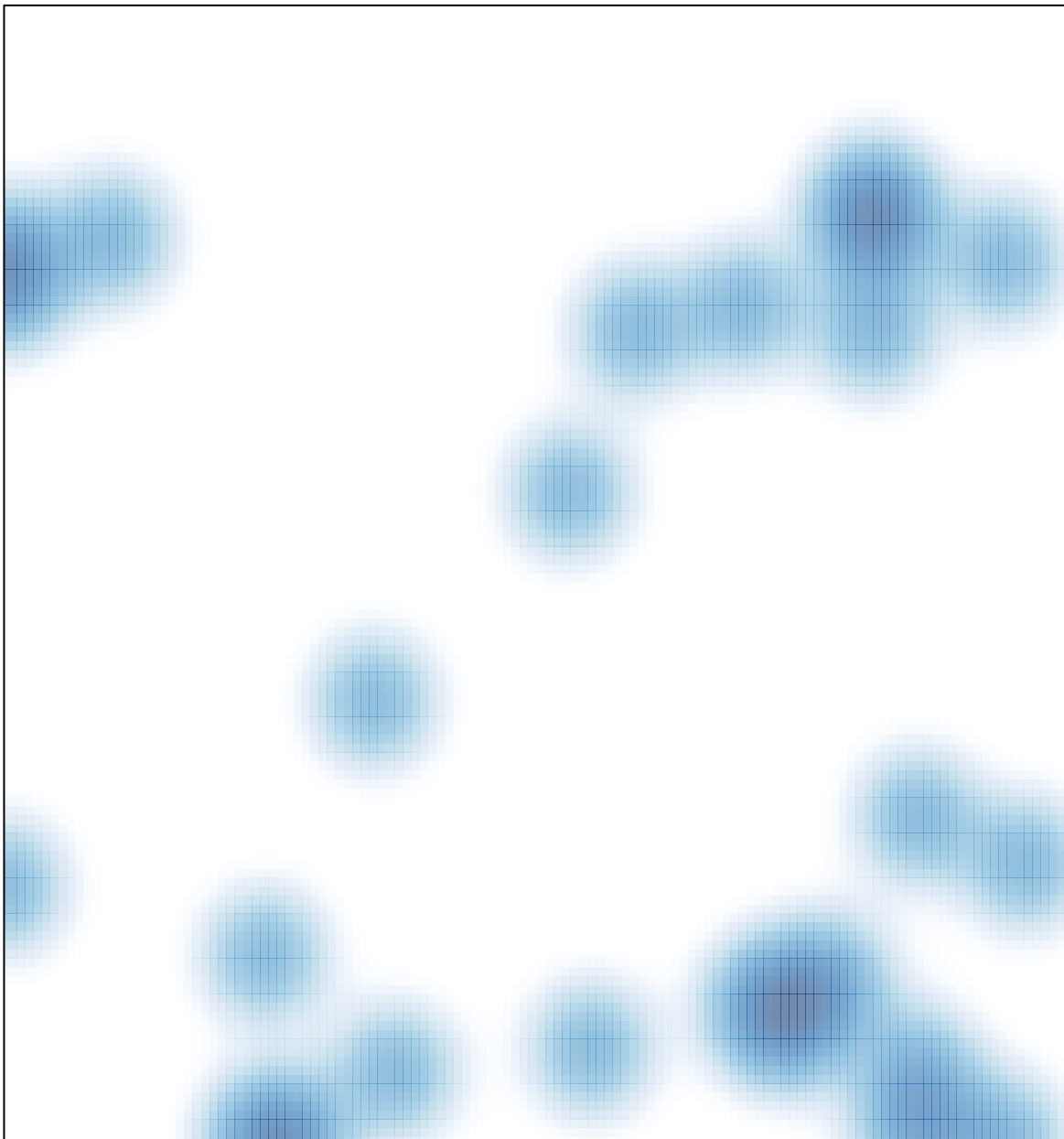


# FIGUEROA\_AML\_METHYLATION\_CLUSTER\_1\_DN



# features = 33  
chi-square p = 0.8

# FIGUEROA\_AML\_METHYLATION\_CLUSTER\_1\_DN



# features = 33 , max = 2