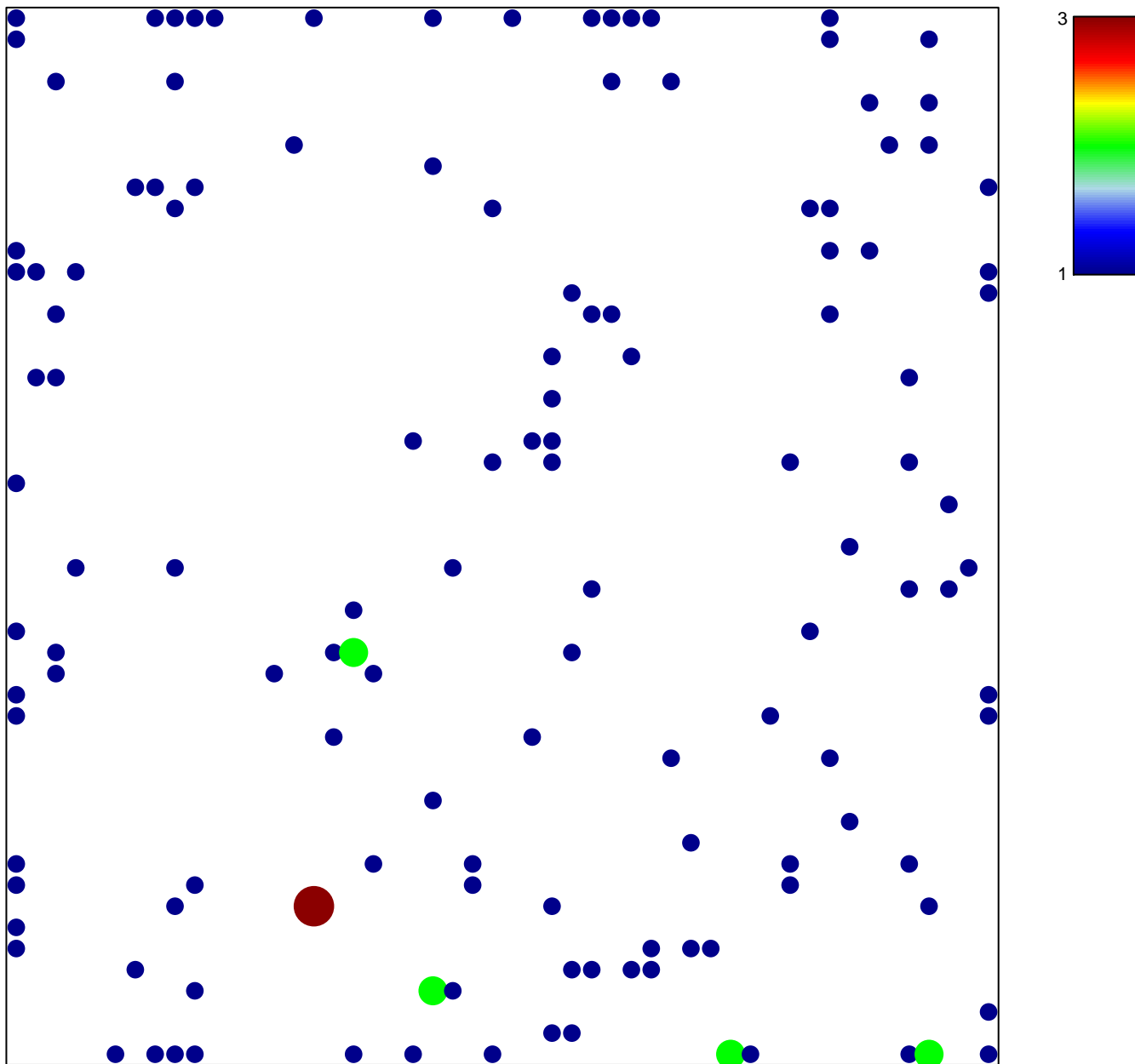
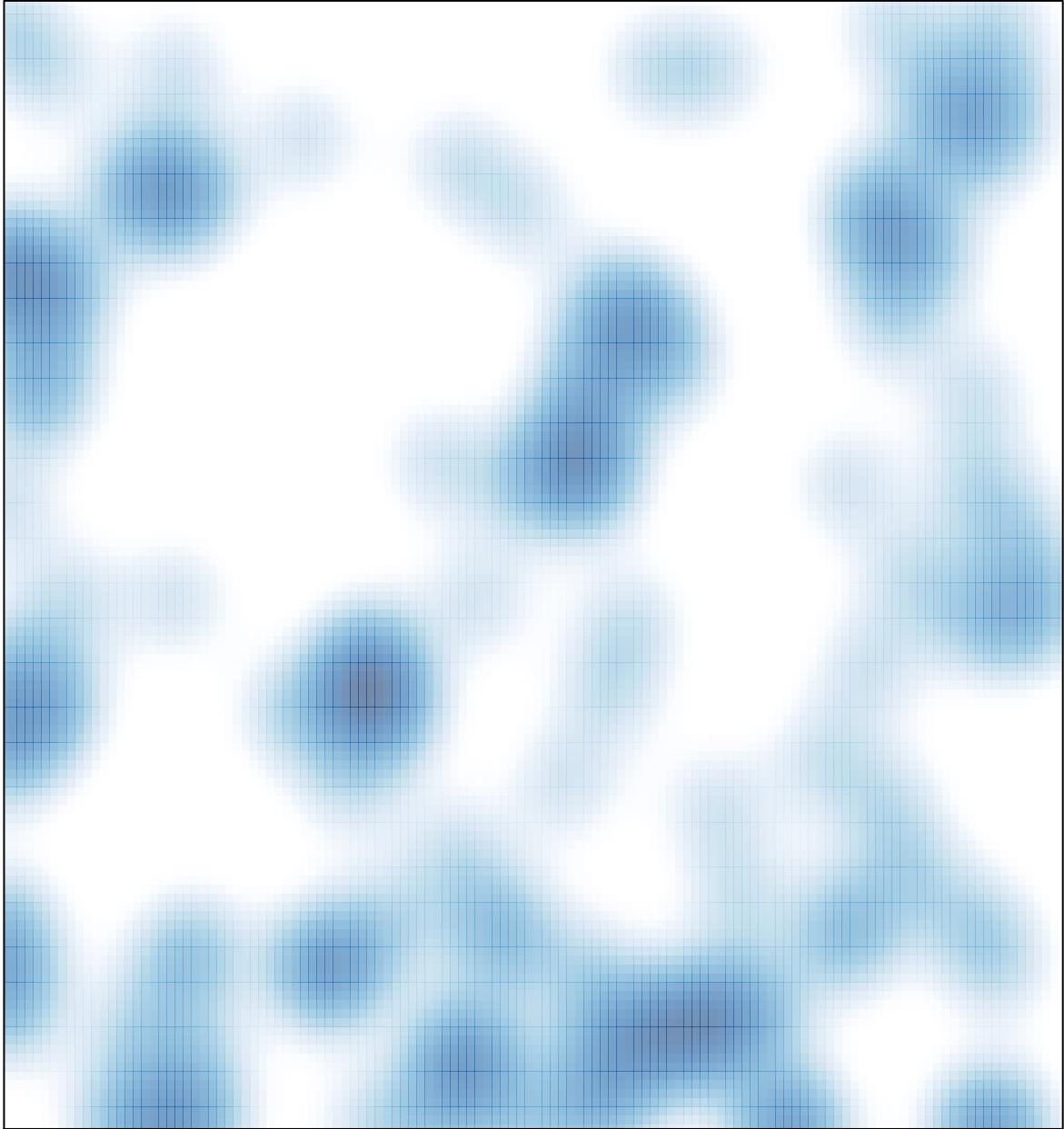


FIGUEROA_AML_METHYLATION_CLUSTER_3_UP



features = 139
chi-square p = 0.77

FIGUEROA_AML_METHYLATION_CLUSTER_3_UP



features = 139 , max = 3