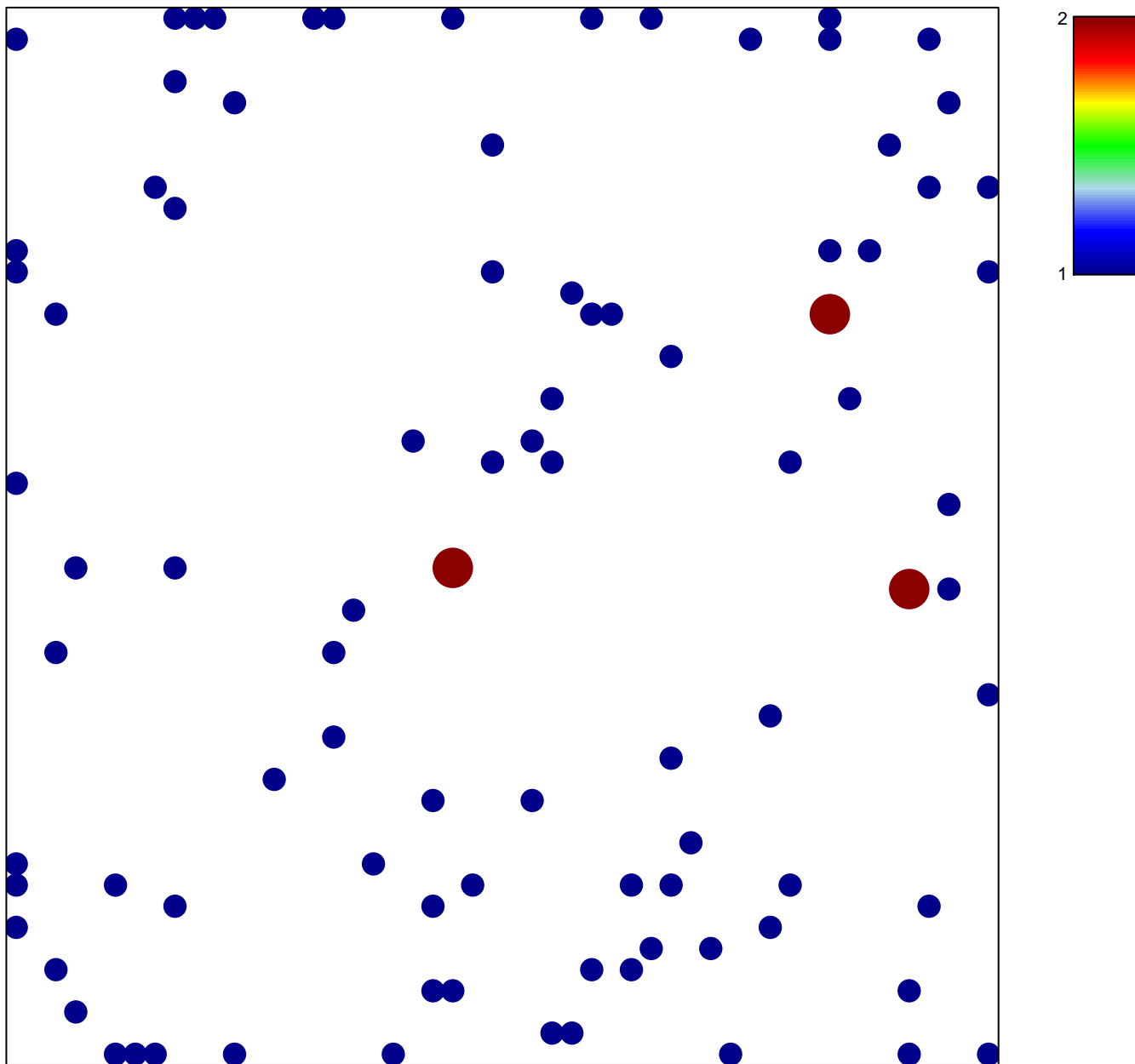
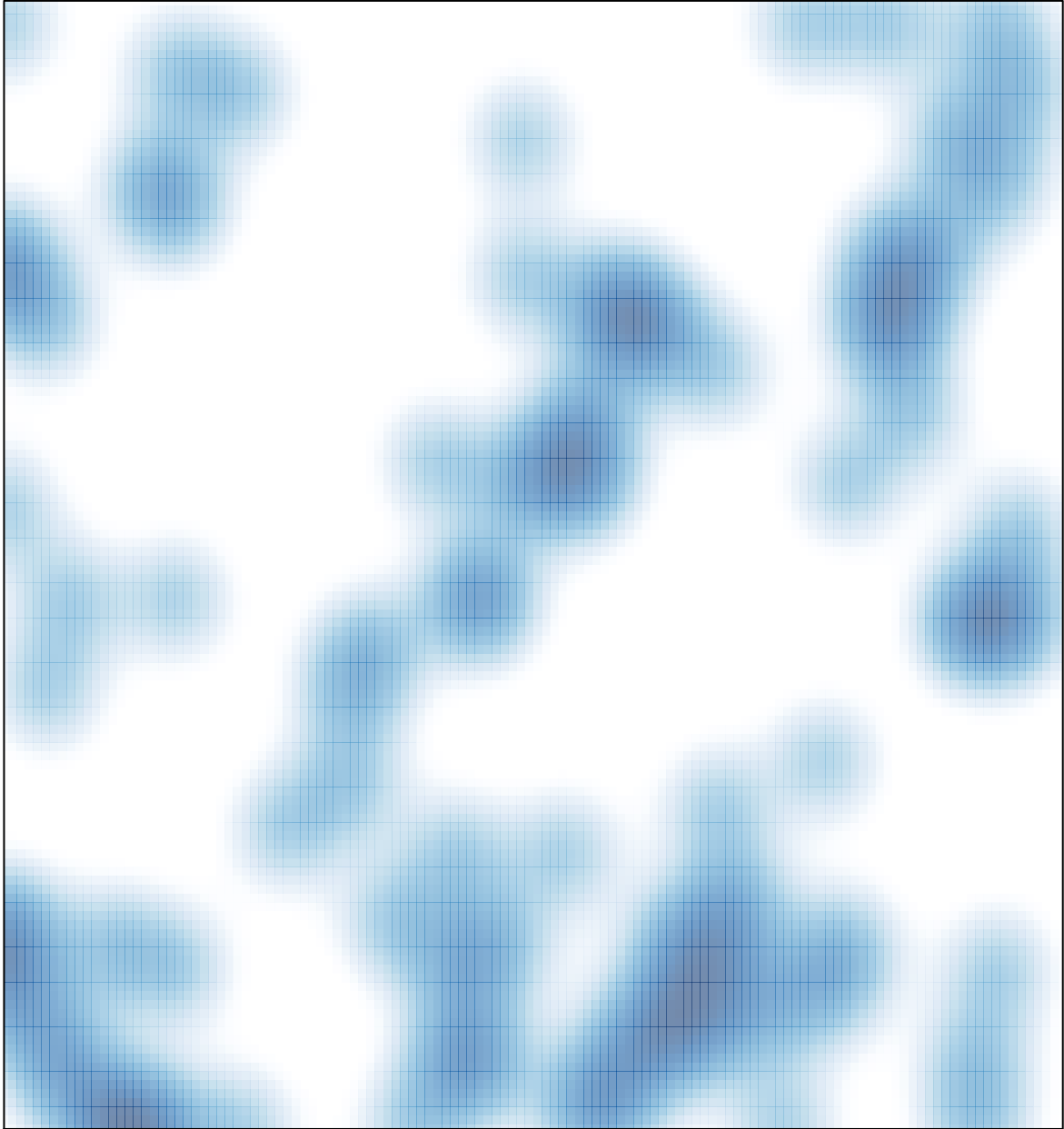


FIGUEROA_AML_METHYLATION_CLUSTER_1_UP



features = 94
chi-square p = 0.8

FIGUEROA_AML_METHYLATION_CLUSTER_1_UP



features = 94 , max = 2