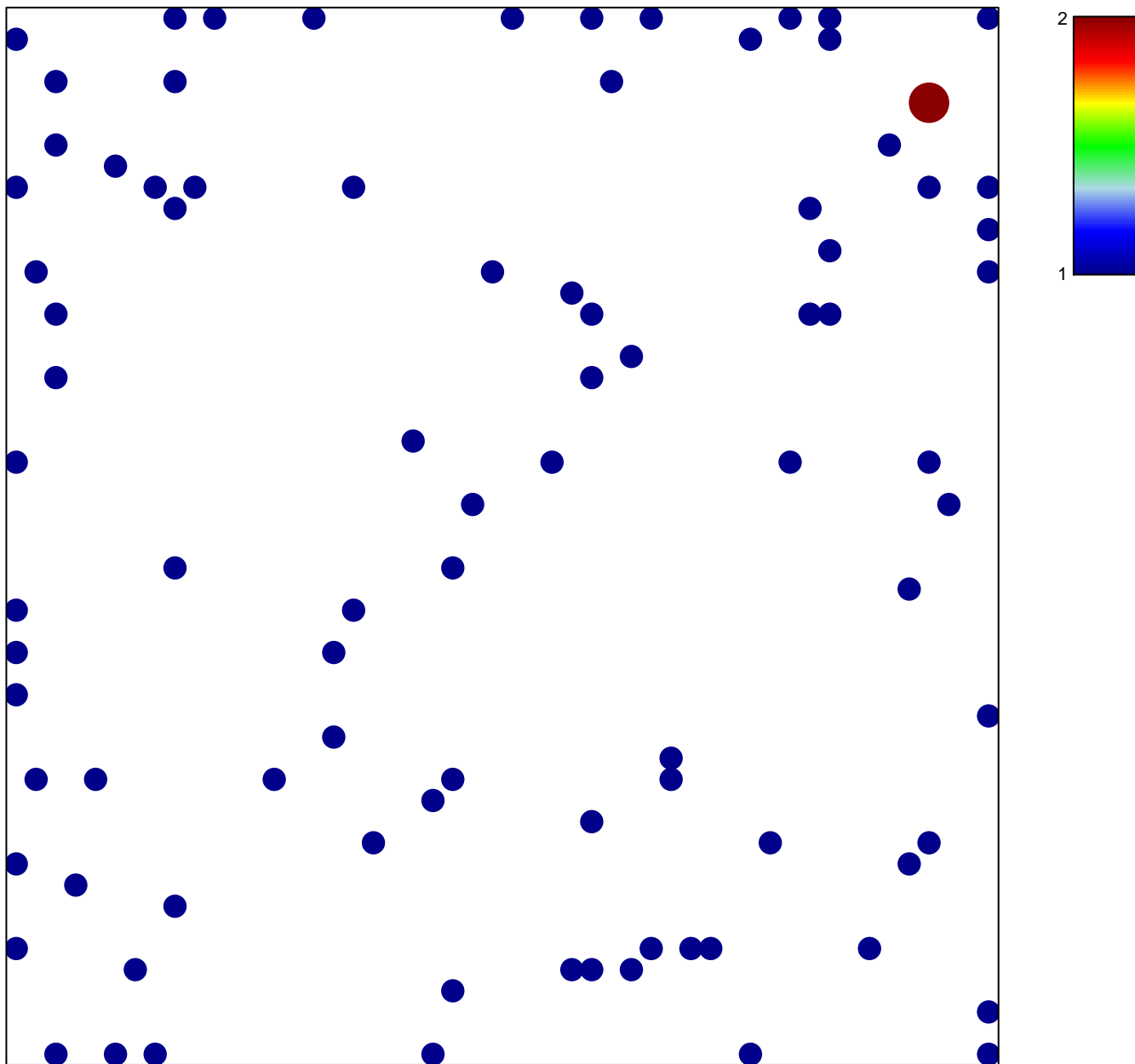
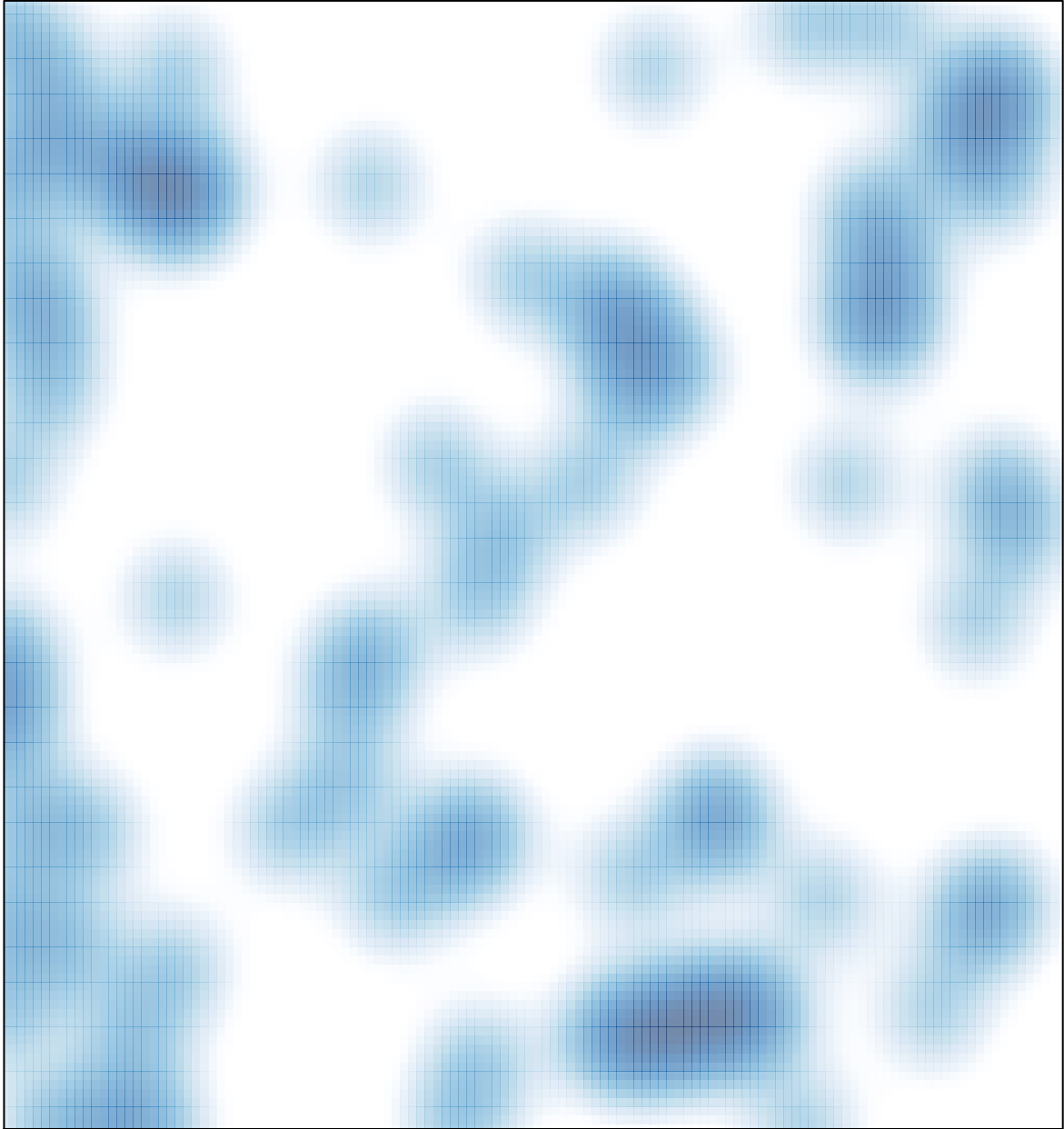


FIGUEROA_AML_METHYLATION_CLUSTER_4_UP



features = 90
chi-square p = 0.79

FIGUEROA_AML_METHYLATION_CLUSTER_4_UP



features = 90 , max = 2