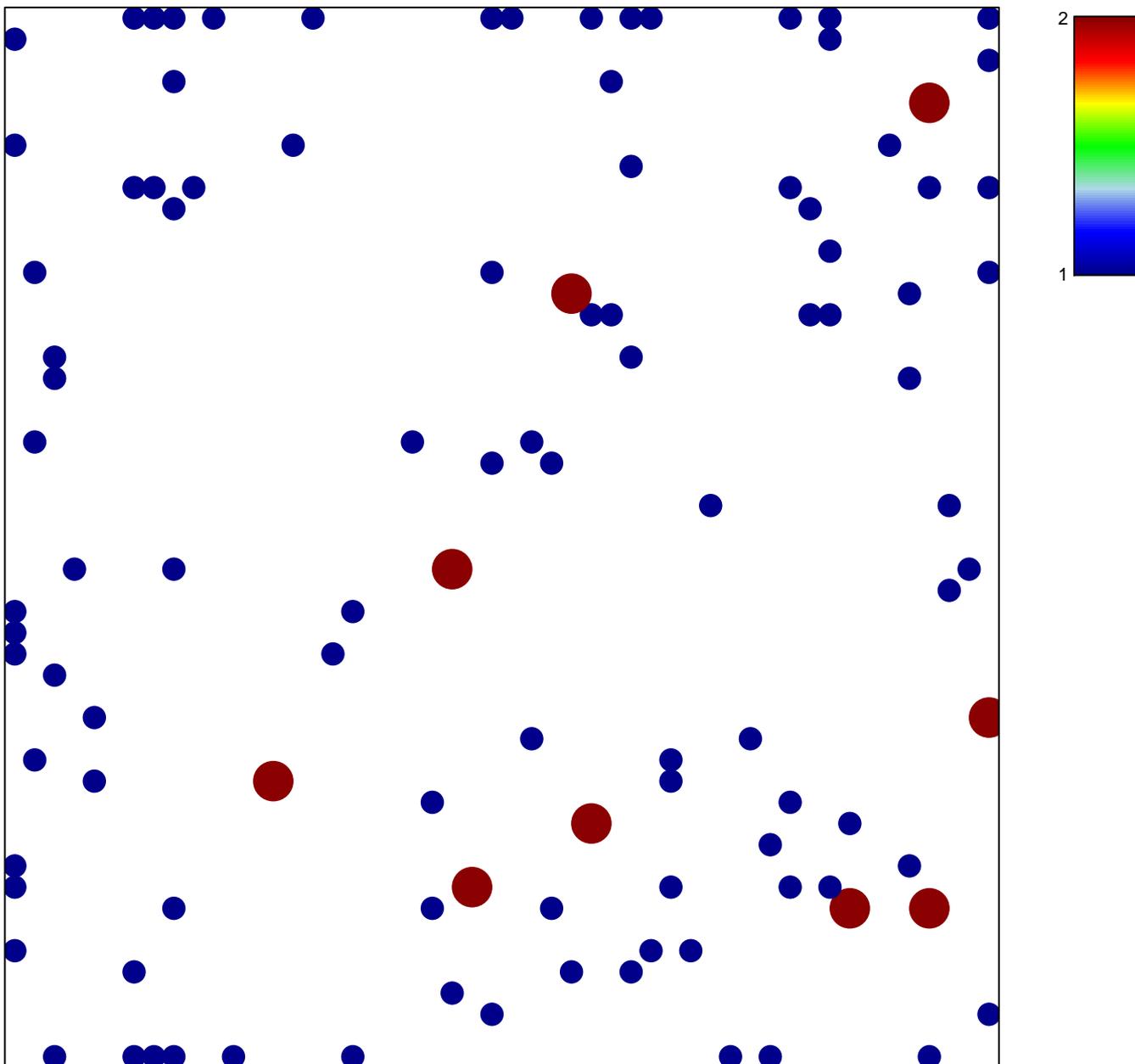
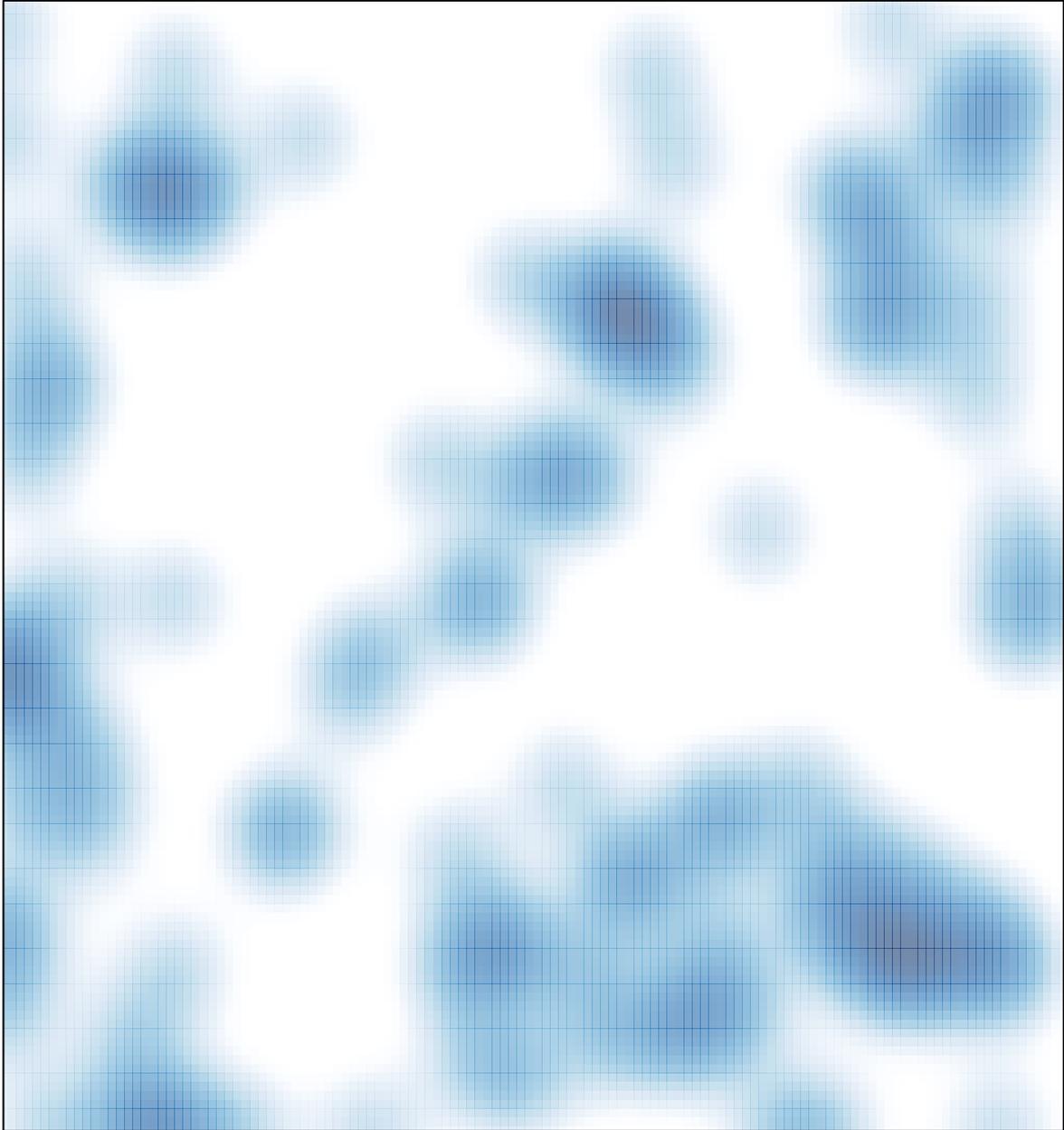


FIGUEROA_AML_METHYLATION_CLUSTER_6_UP



features = 116
chi-square p = 0.78

FIGUEROA_AML_METHYLATION_CLUSTER_6_UP



features = 116 , max = 2