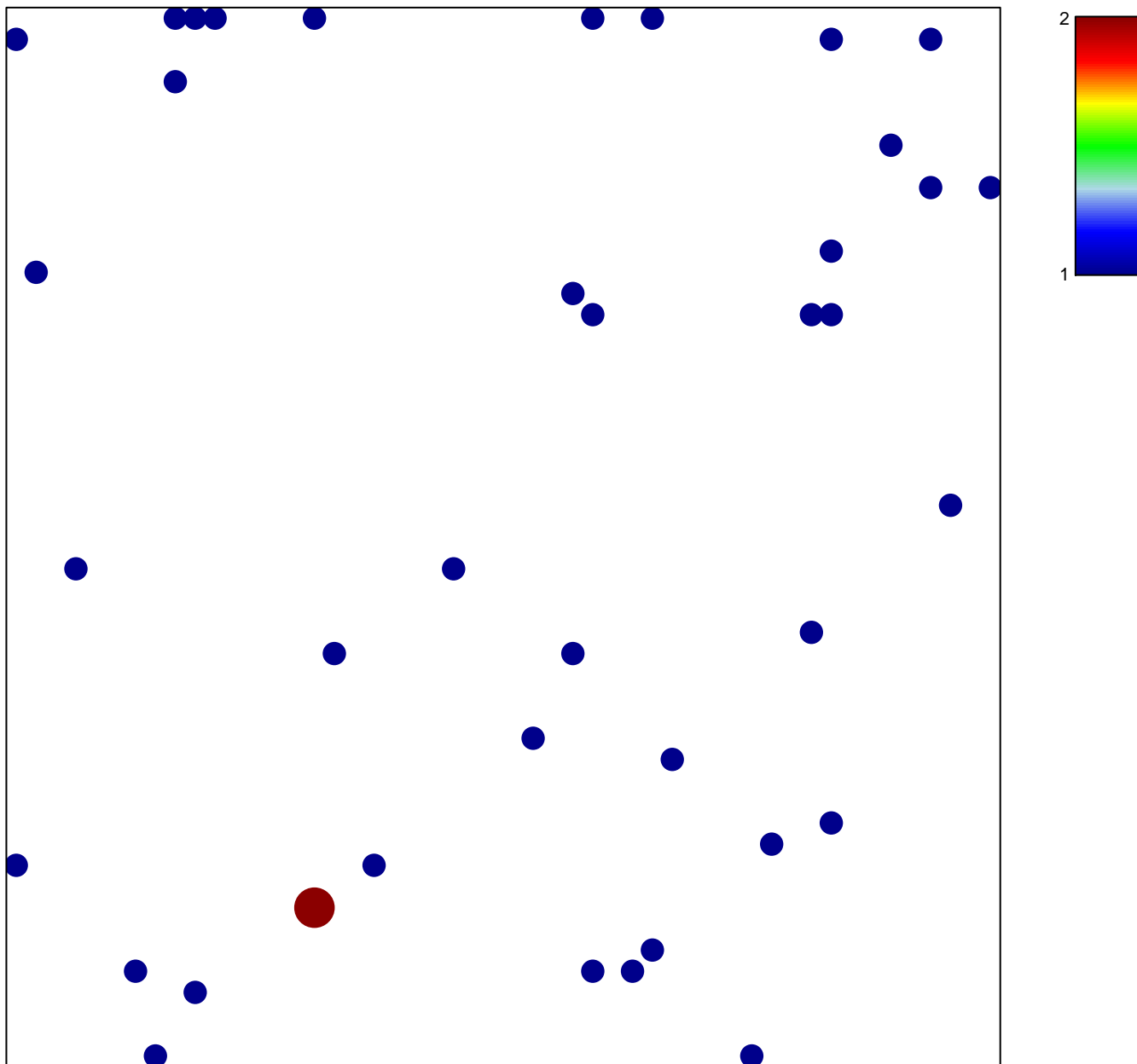


FIGUEROA_AML_METHYLATION_CLUSTER_2_UP



features = 40
chi-square p = 0.81

FIGUEROA_AML_METHYLATION_CLUSTER_2_UP



features = 40 , max = 2