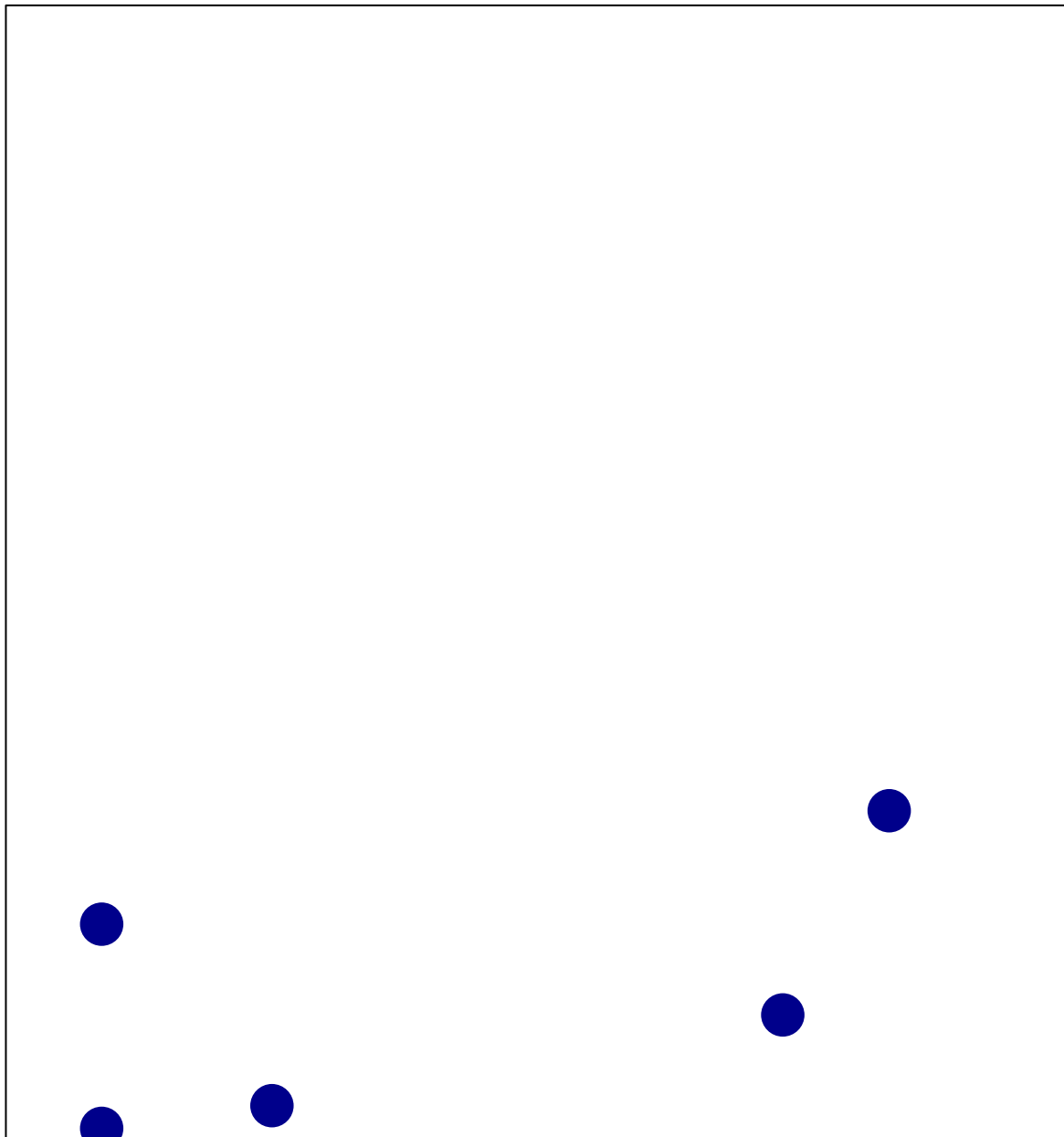


FIGUEROA_AML_METHYLATION_CLUSTER_2_DN



features = 5
chi-square p = 0.83

FIGUEROA_AML_METHYLATION_CLUSTER_2_DN



features = 5 , max = 1