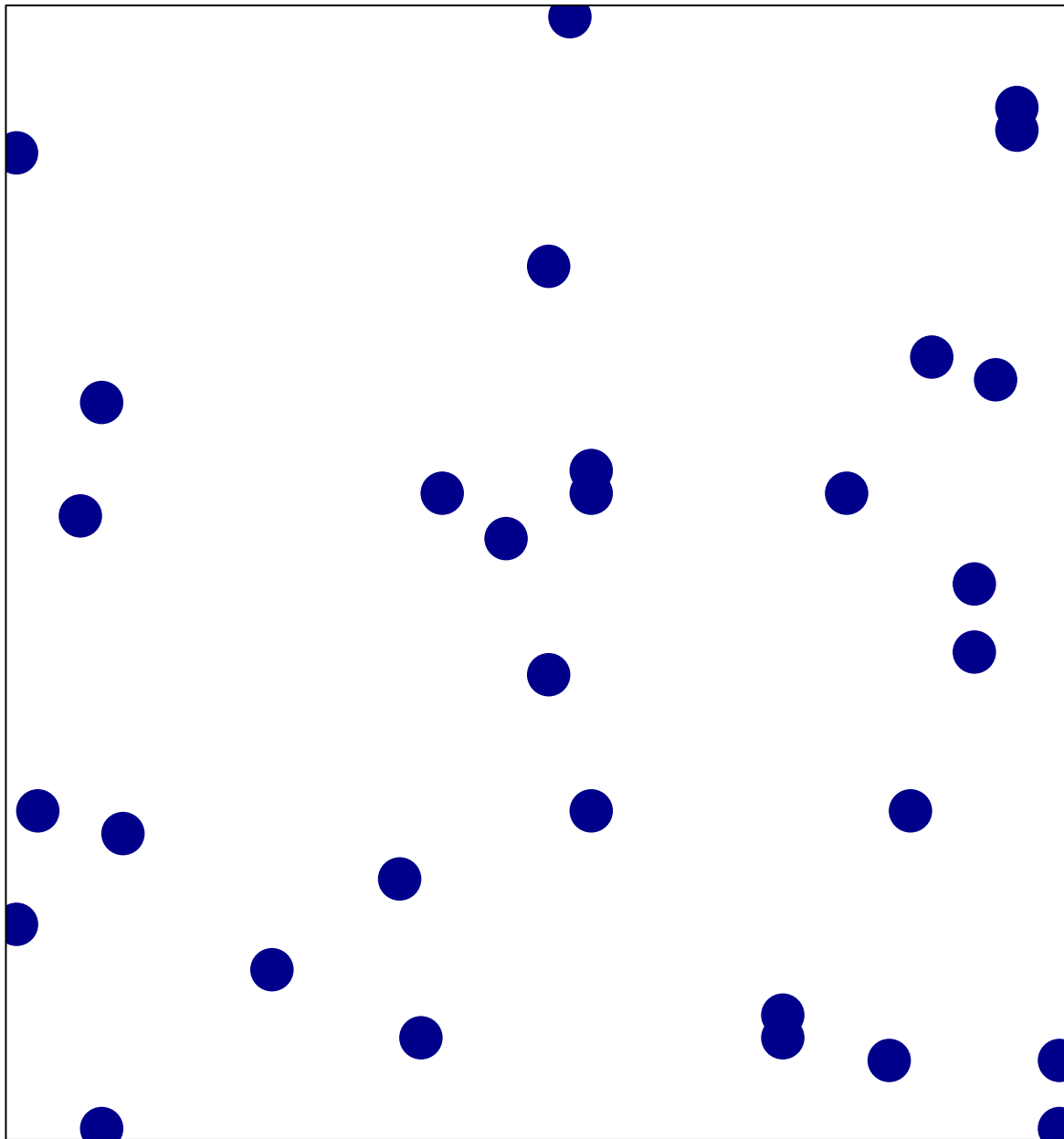
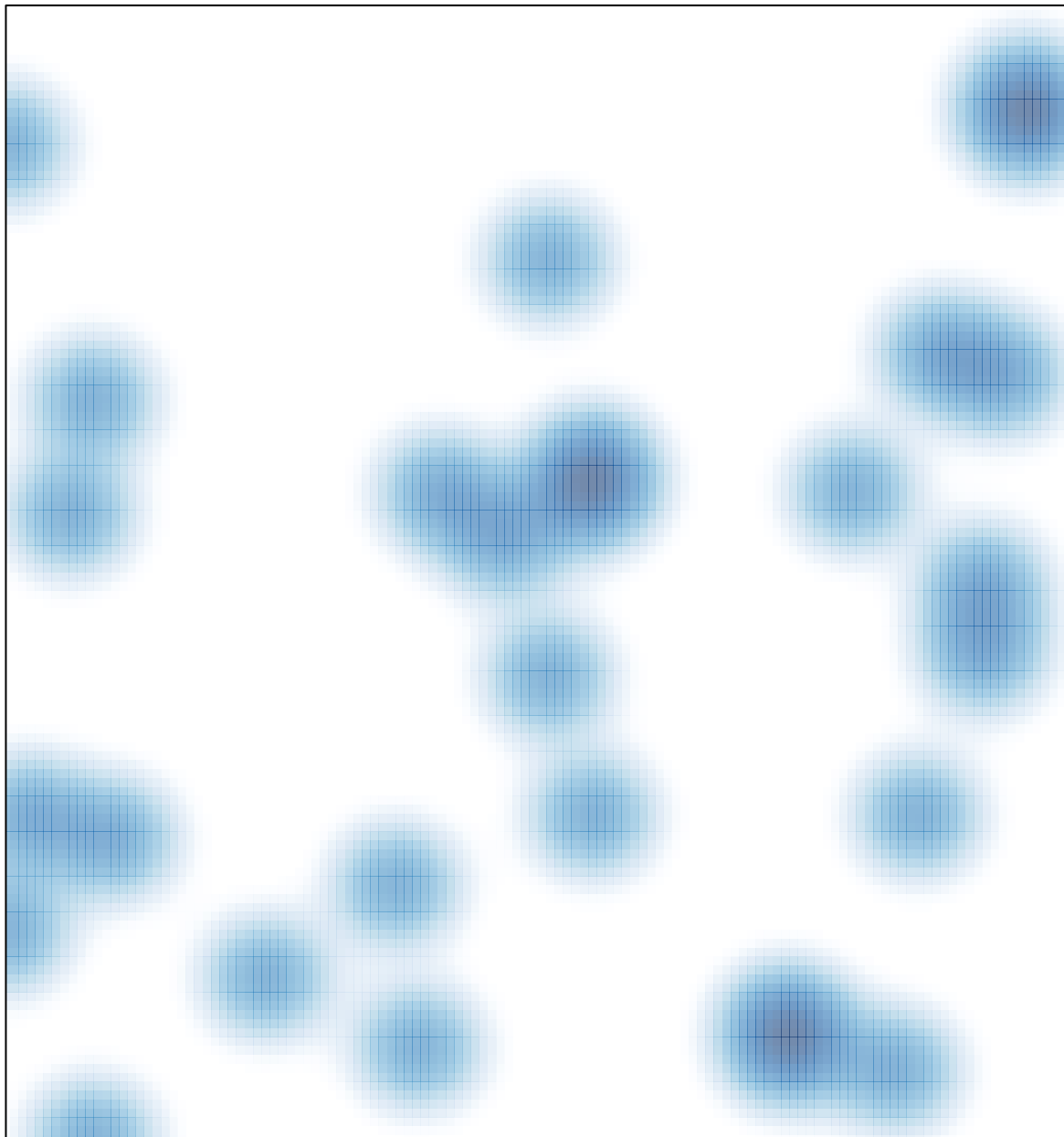


FIGUEROA_AML_METHYLATION_CLUSTER_5_DN



features = 31
chi-square p = 0.84

FIGUEROA_AML_METHYLATION_CLUSTER_5_DN



features = 31 , max = 1