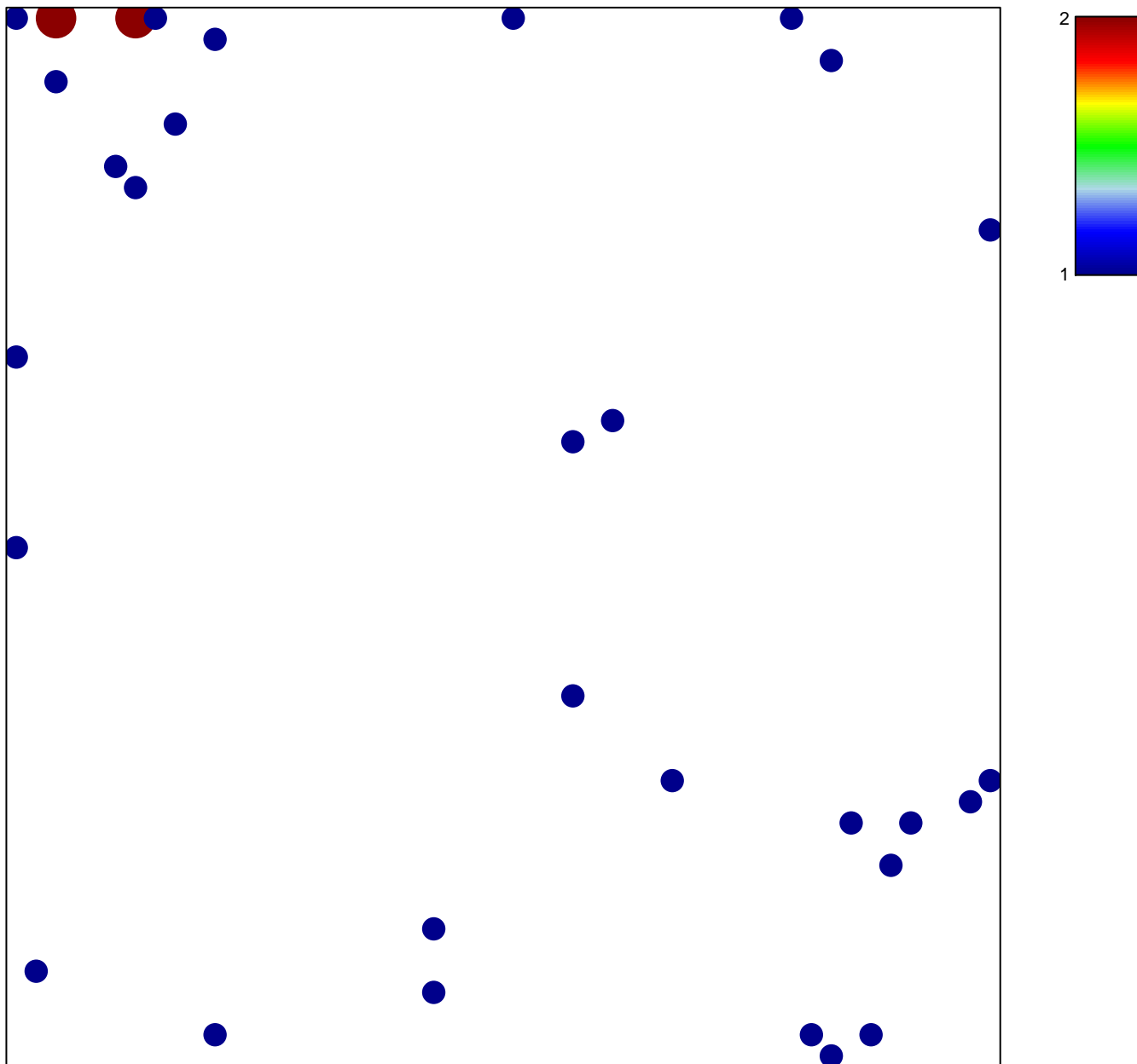
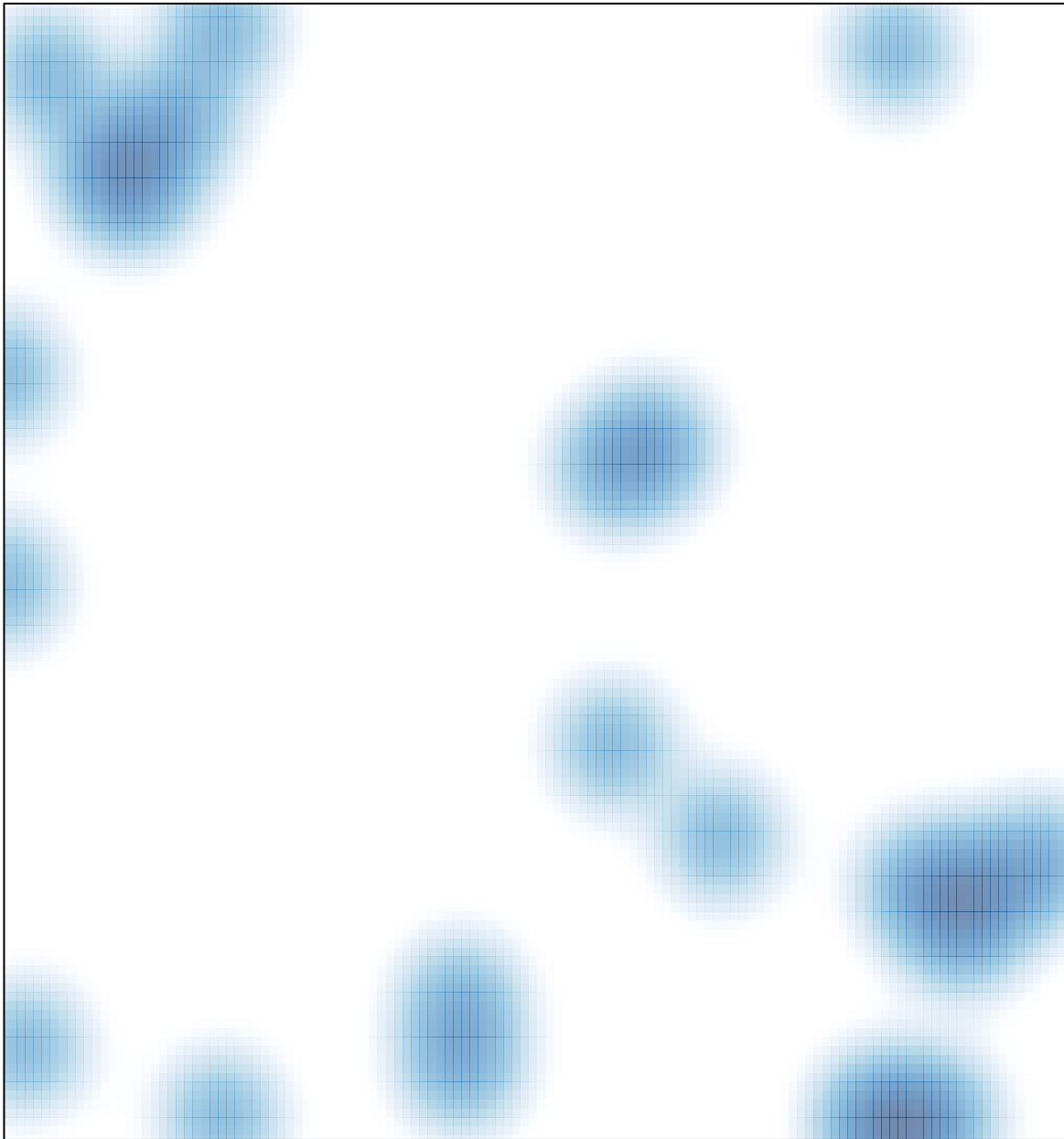


BOYLAN_MULTIPLE_MYELOMA_C_CLUSTER_UP



features = 33
chi-square p = 0.79

BOYLAN_MULTIPLE_MYELOMA_C_CLUSTER_UP



features = 33 , max = 2