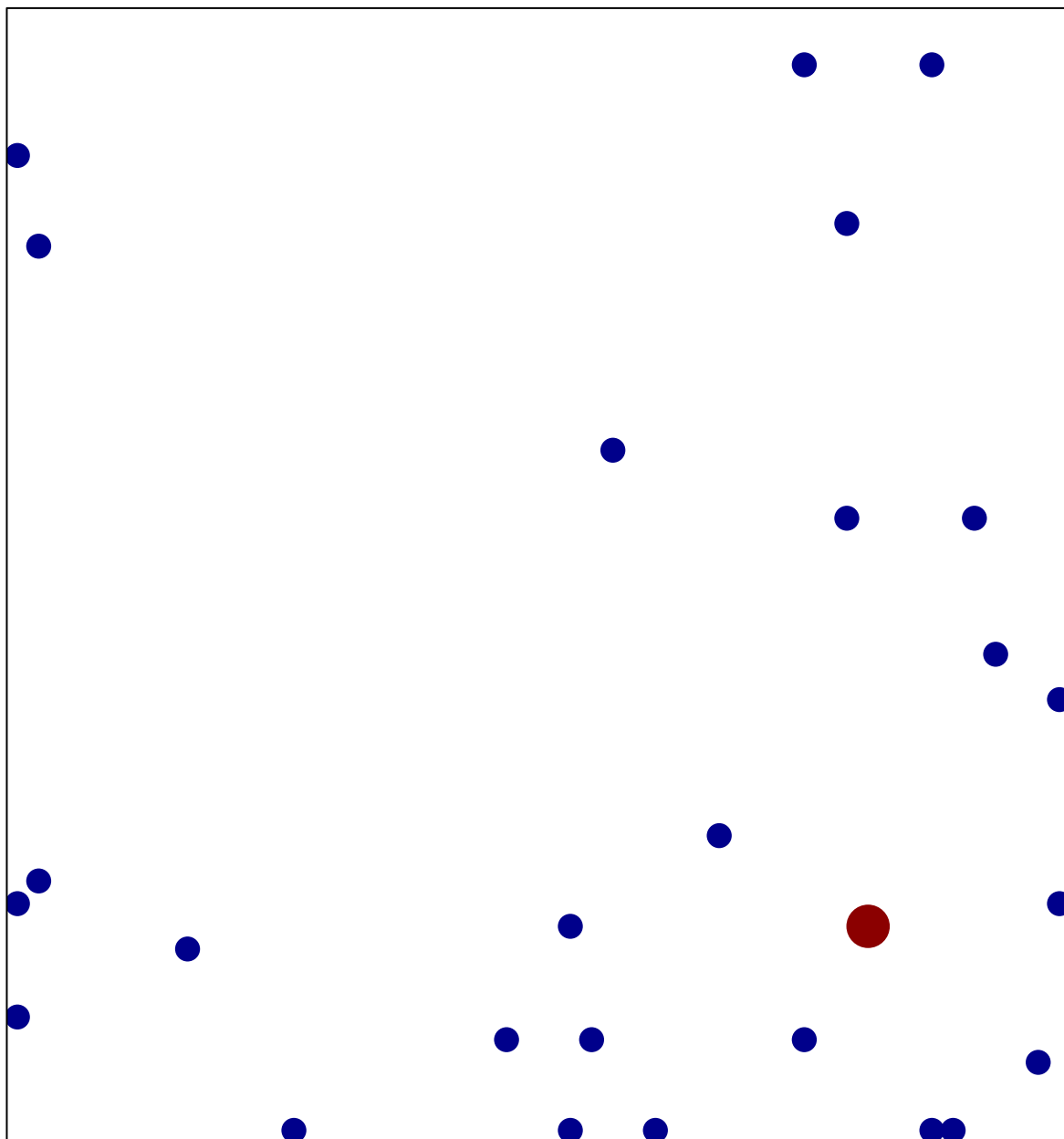
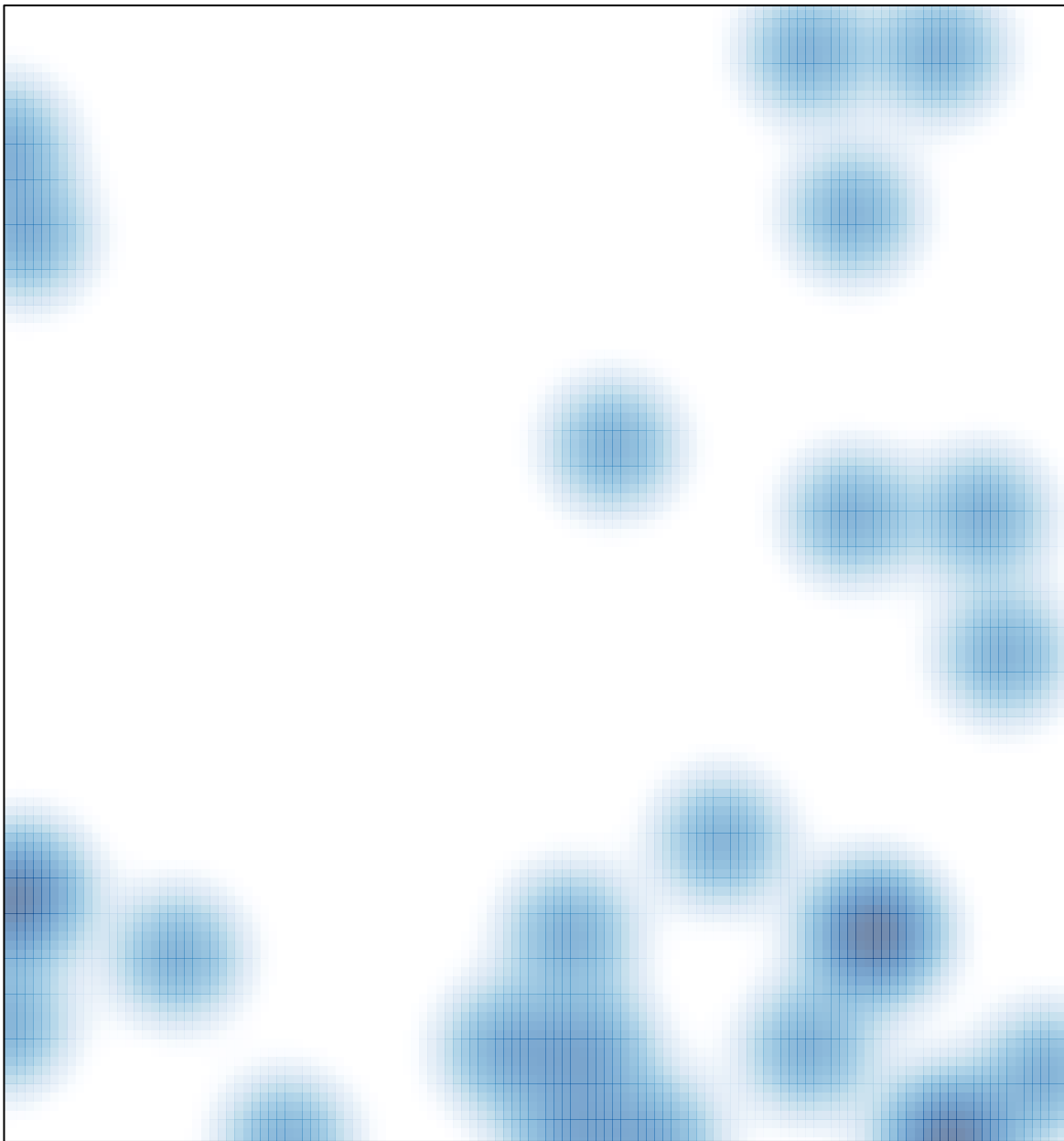


BOYLAN_MULTIPLE_MYELOMA_C_CLUSTER_DN



features = 28
chi-square p = 0.81

BOYLAN_MULTIPLE_MYELOMA_C_CLUSTER_DN



features = 28 , max = 2