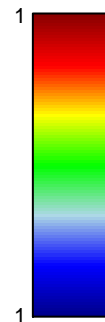
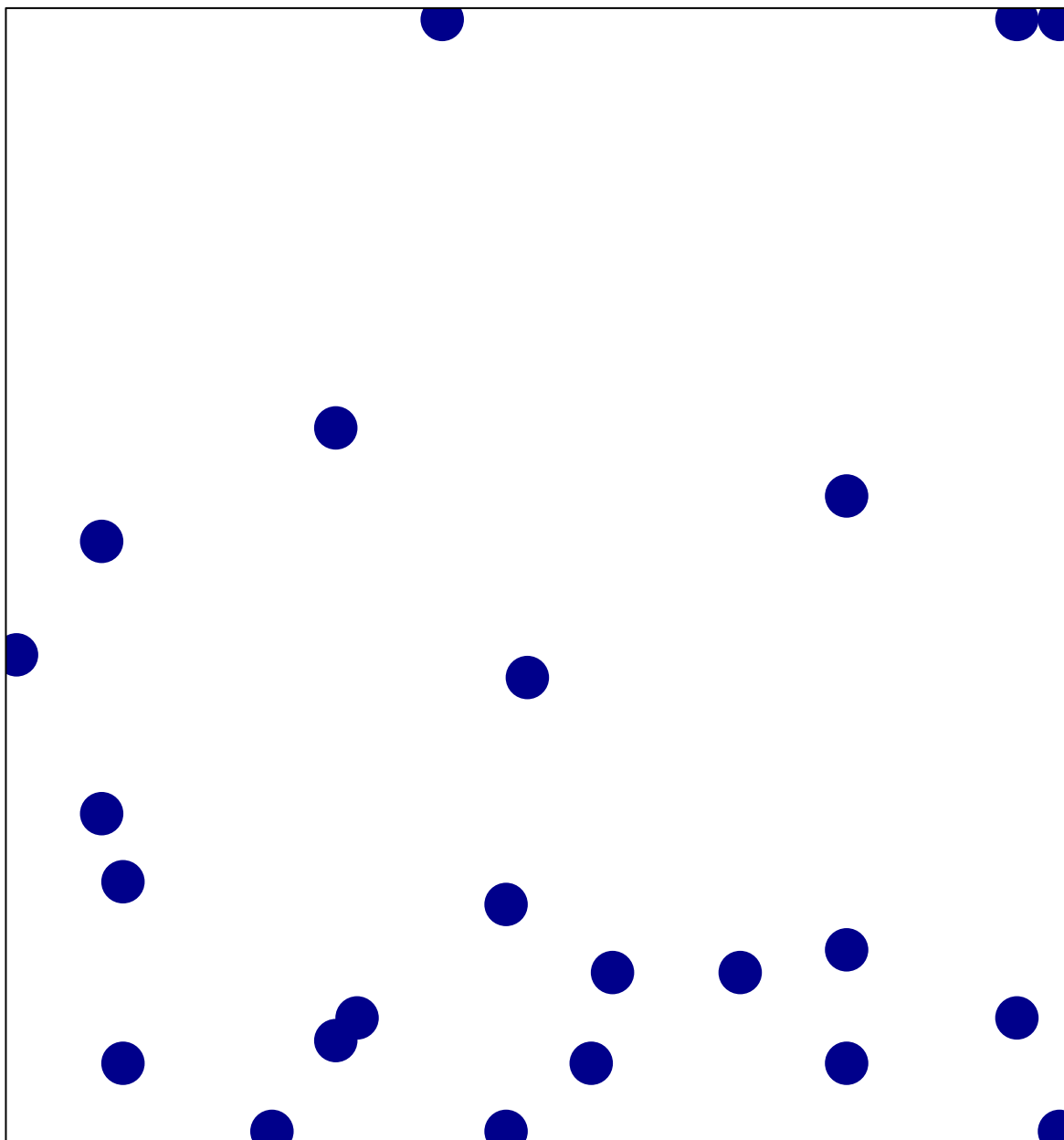


BOYLAN_MULTIPLE_MYELOMA_D_CLUSTER_UP



features = 23
chi-square p = 0.81

BOYLAN_MULTIPLE_MYELOMA_D_CLUSTER_UP



features = 23 , max = 1