DAWSON_METHYLATED_IN_LYMPHOMA_TCL1
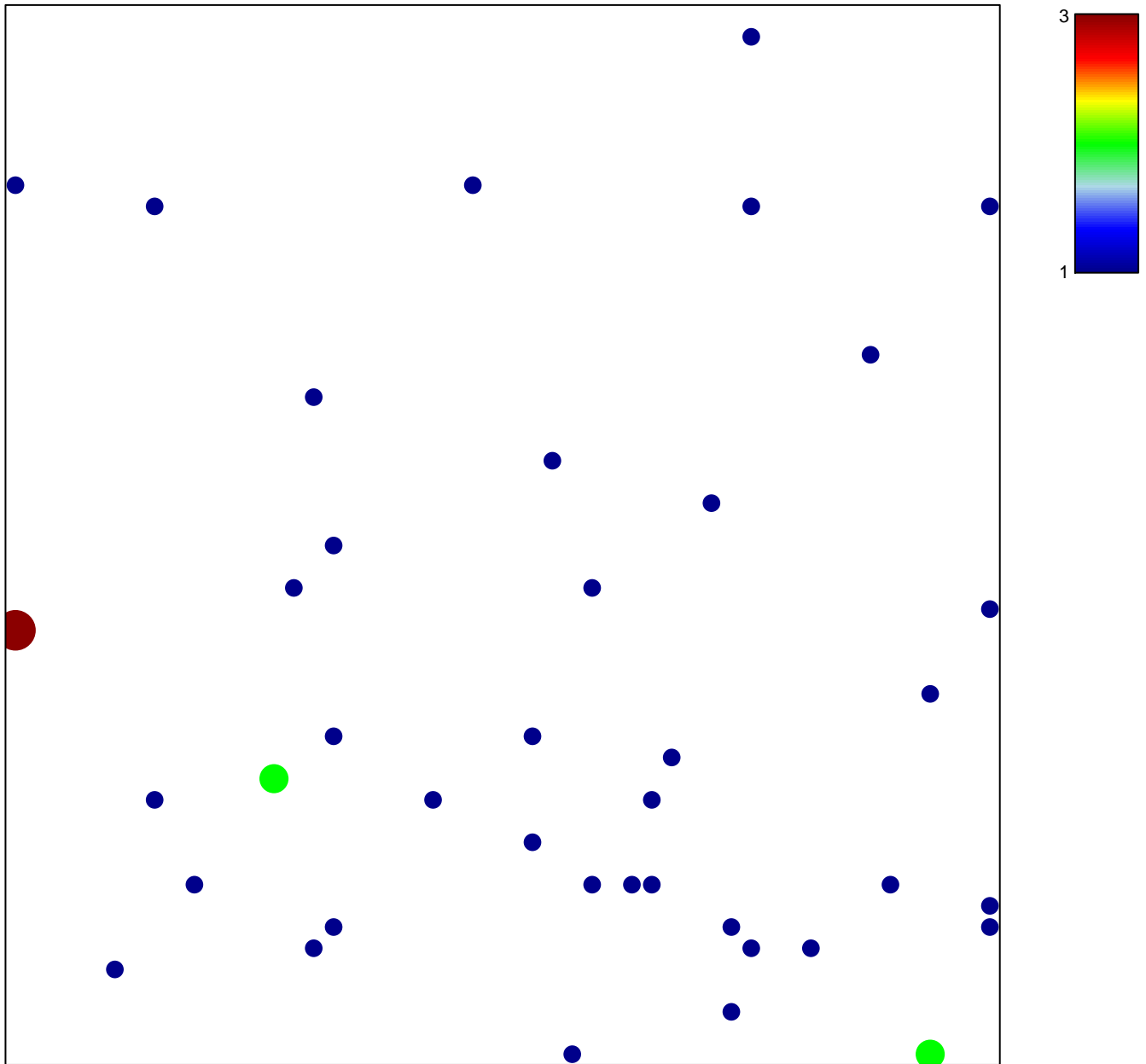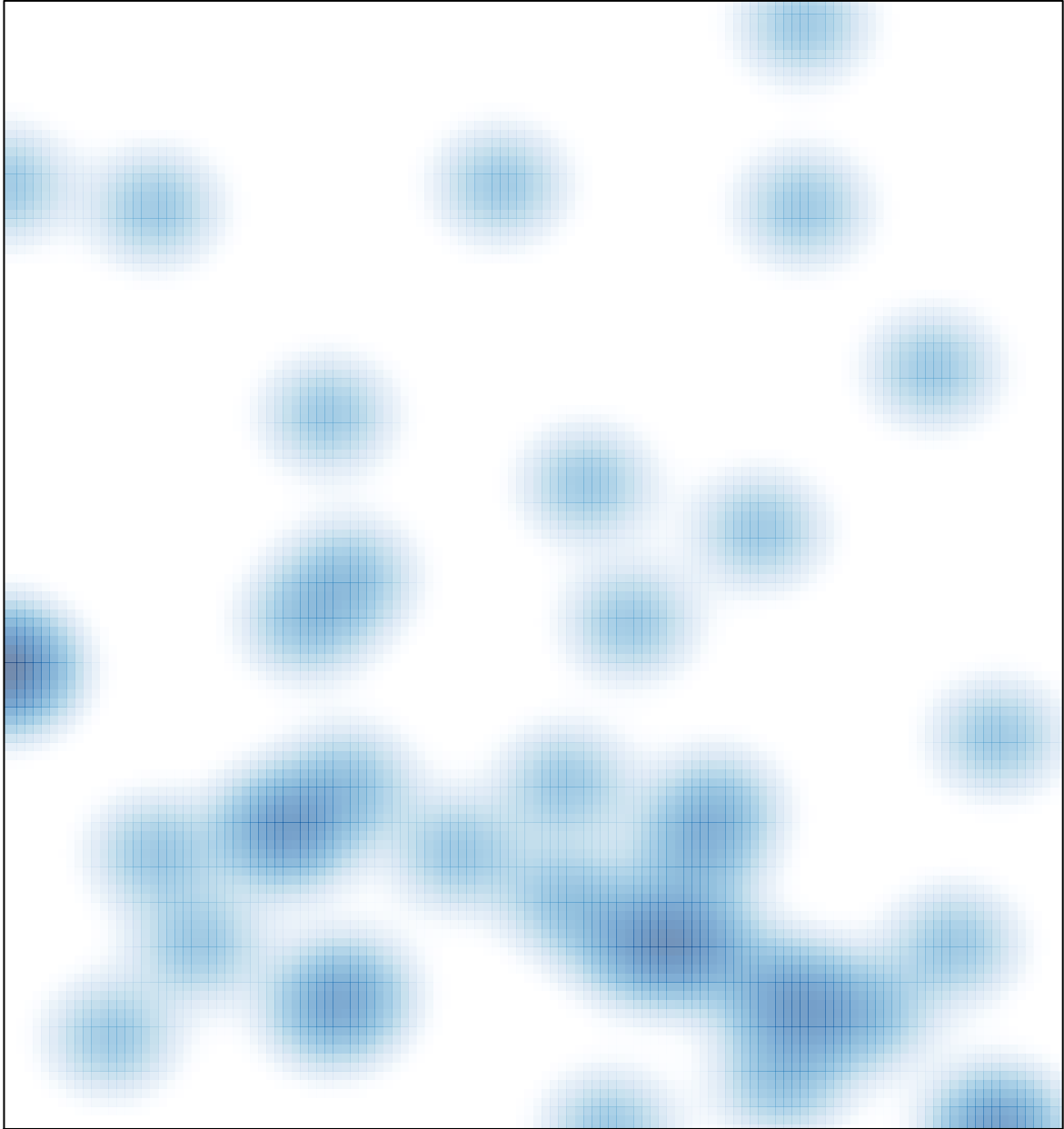

features = 44

chi−square p = 0.81

DAWSON_METHYLATED_IN_LYMPHOMA_TCL1

features = 44 , max = 3