FAELT_B_CLL_WITH_VH_REARRANGEMENTS_DN
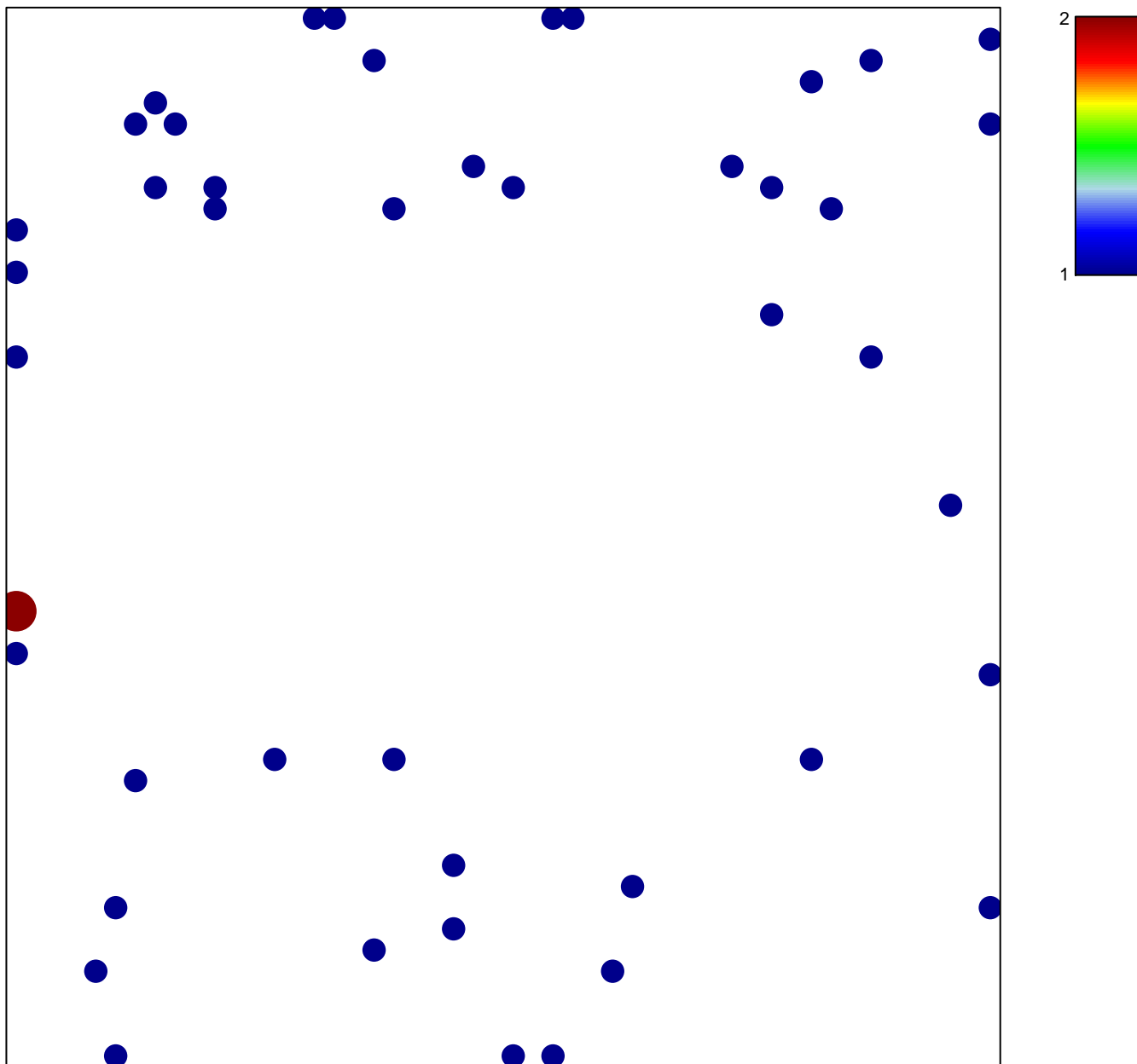

features = 46
chi−square p = 0.81

FAELT_B_CLL_WITH_VH_REARRANGEMENTS_DN

features = 46 , max = 2