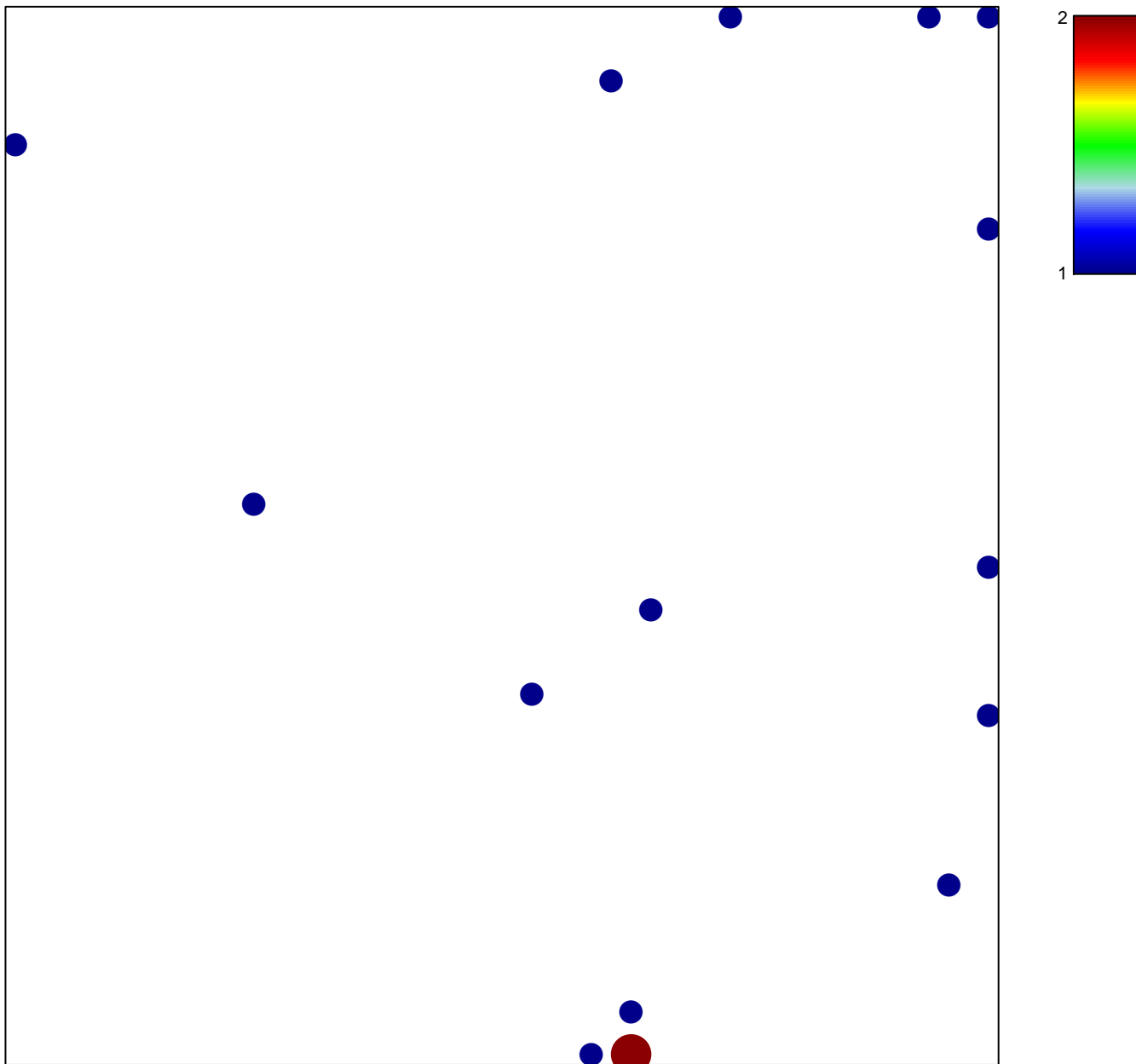
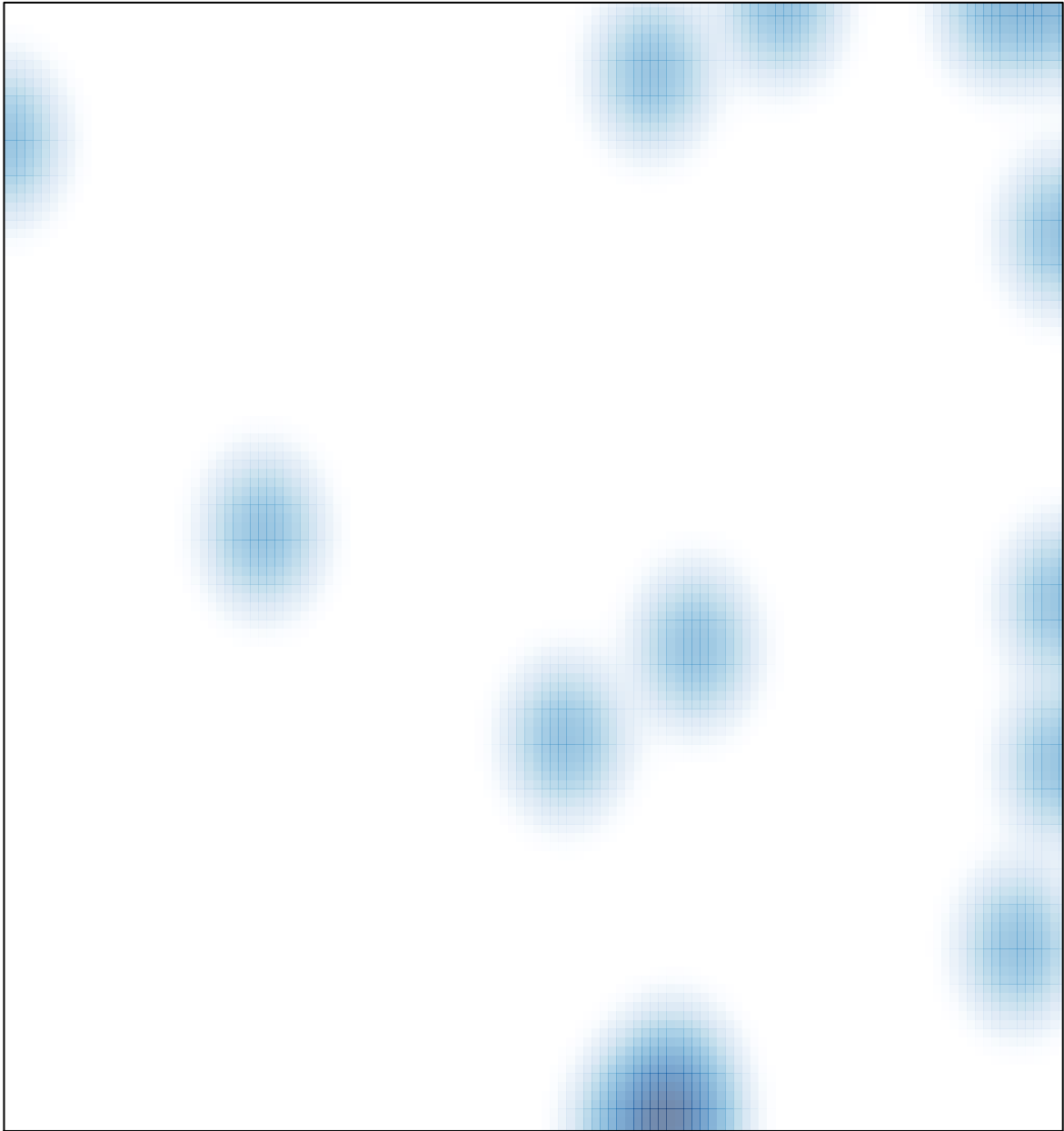


SENGUPTA_NASOPHARYNGEAL_CARCINOMA_WITH_LMP1_UP



features = 16
chi-square p = 0.8

UPTA_NASOPHARYNGEAL_CARCINOMA_WITH_LM



features = 16 , max = 2