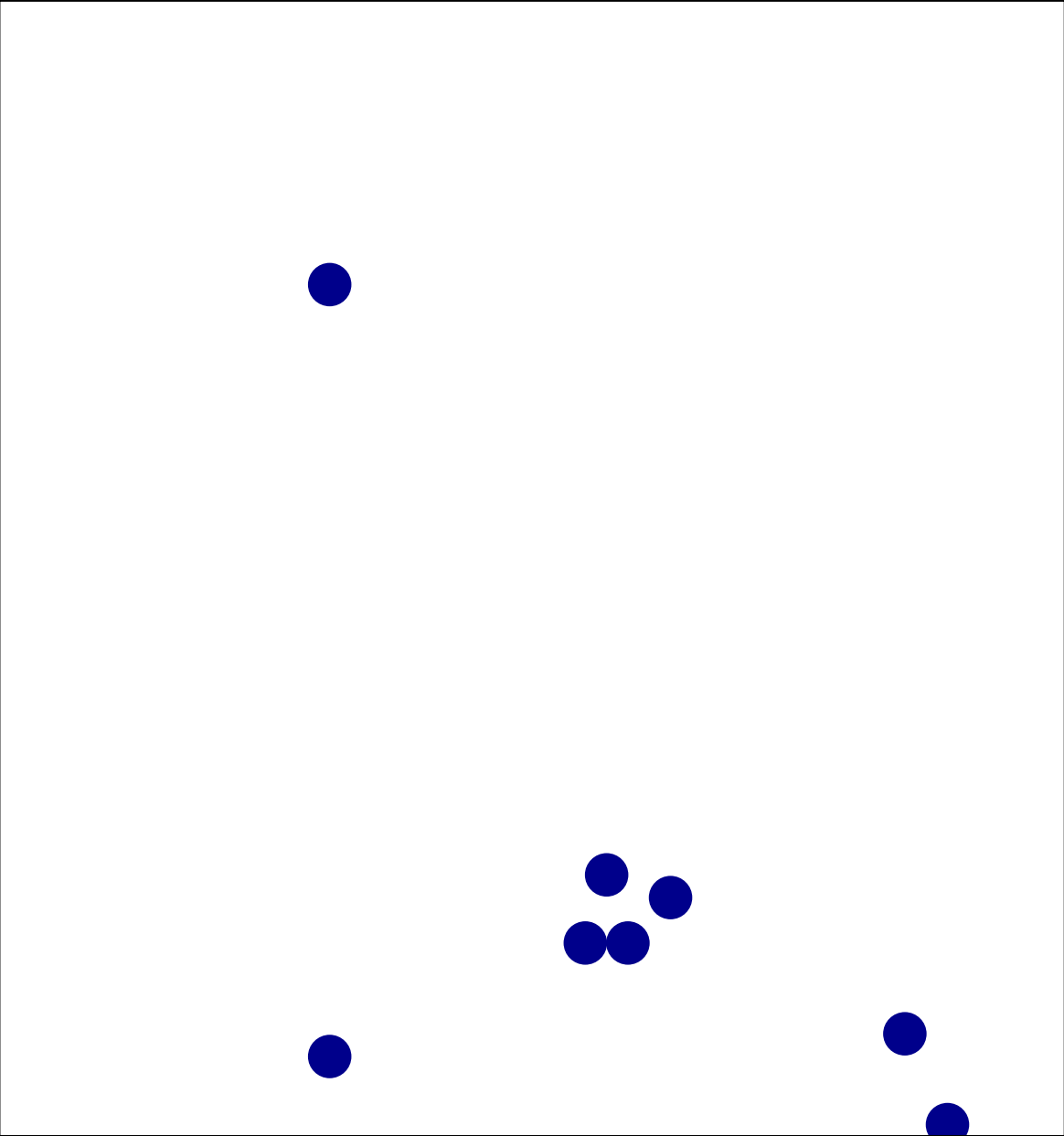
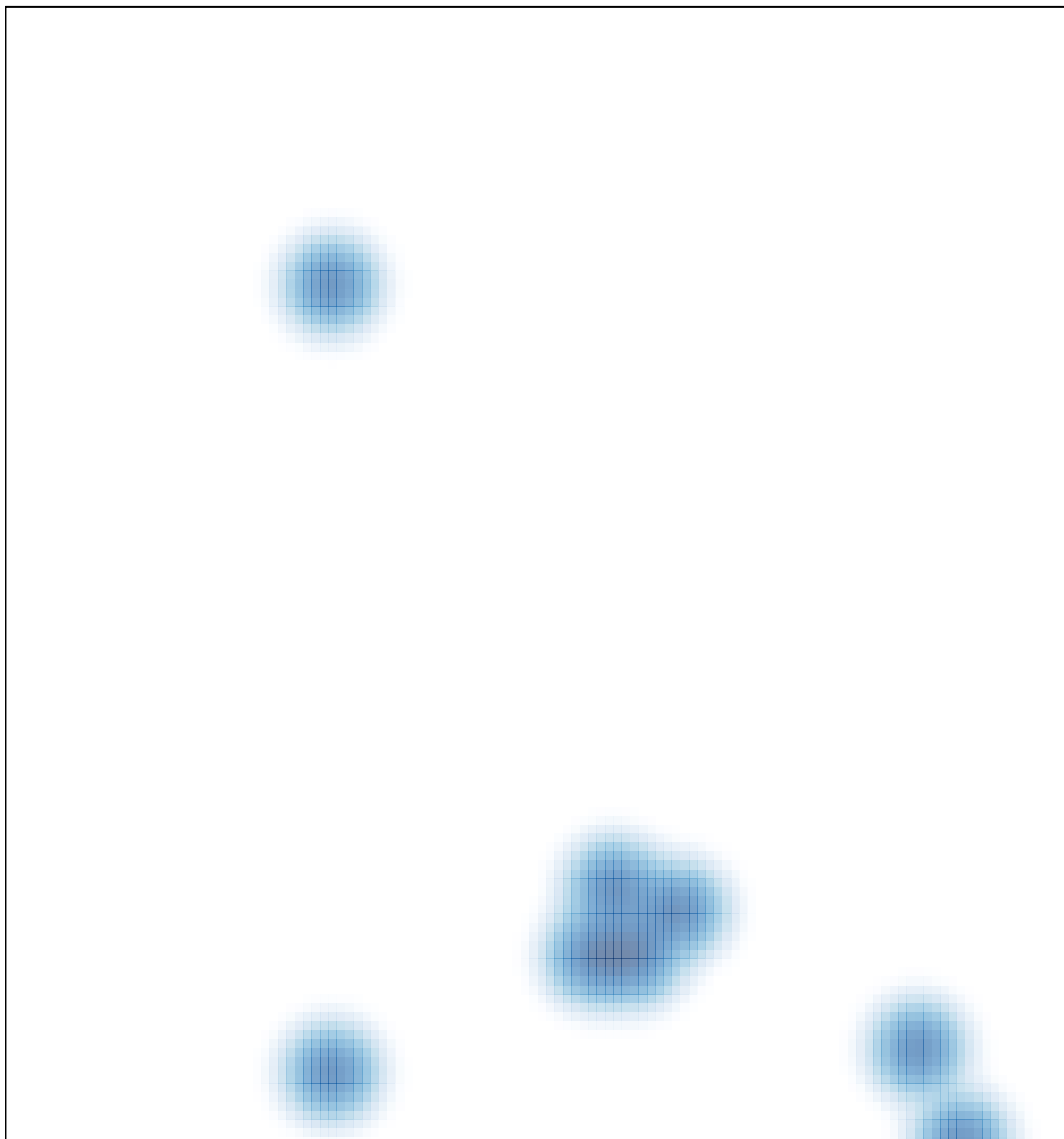


TCGA_GLIOMASTOMA_MUTATED



features = 8
chi-square p = 0.83

TCGA_GLIOBLASTOMA_MUTATED



features = 8 , max = 1