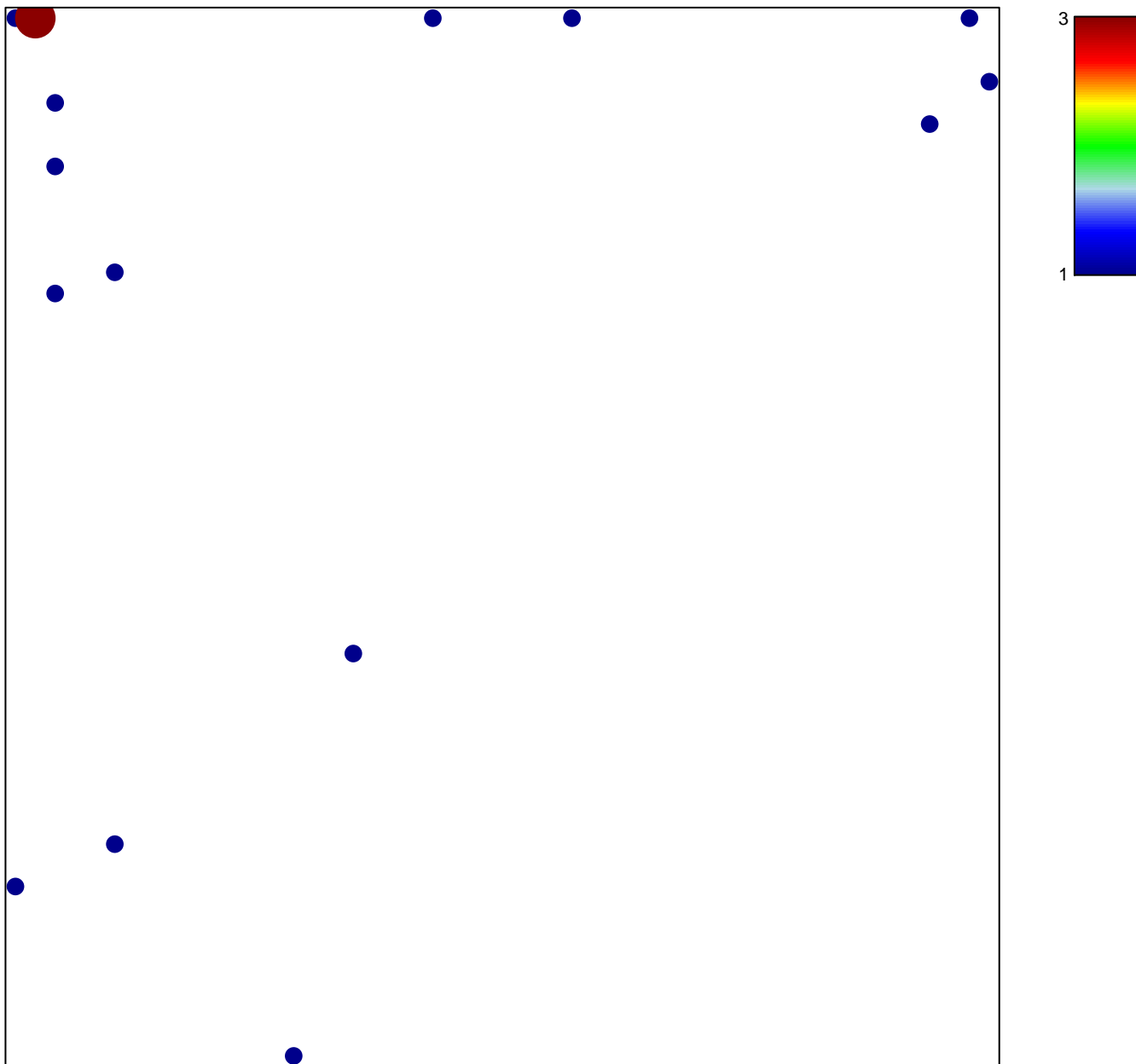


E_RESOLUTION_OF_AP_SITES_VIA_THE_MULTIPLE_NUCLEOTIDE_PATCH_REPLACEME



features = 17
chi-square p = 0.76

FAP_SITES_VIA_THE_MULTIPLE_NUCLEOTIDE_PAT



features = 17 , max = 3