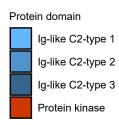
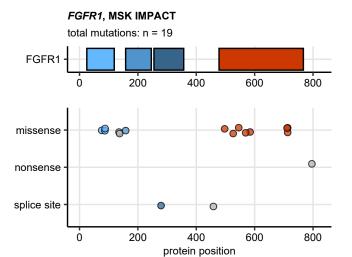
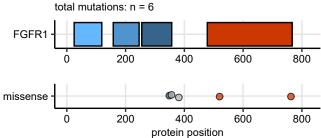
FGFR1, GENIE BLCA total mutations: n = 66 FGFR1 missense nonsense splice region splice site

protein position







FGFR1, TCGA BLCA