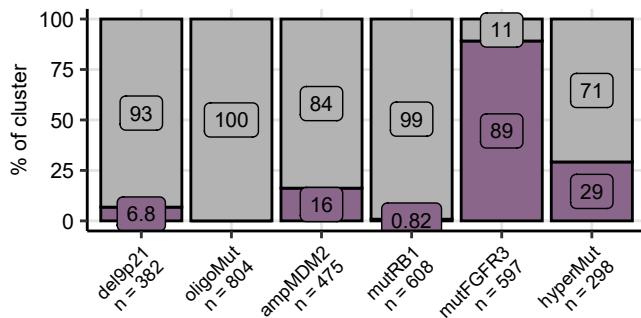
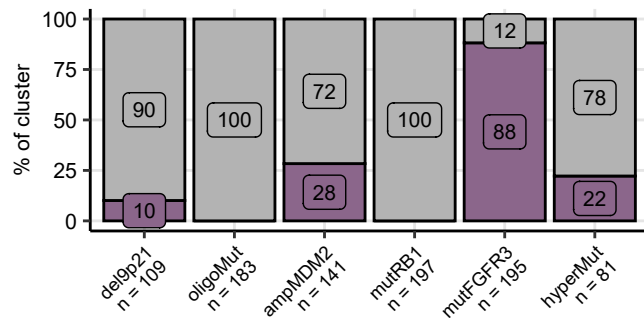


A***FGFR3* mutation, GENIE BLCA**

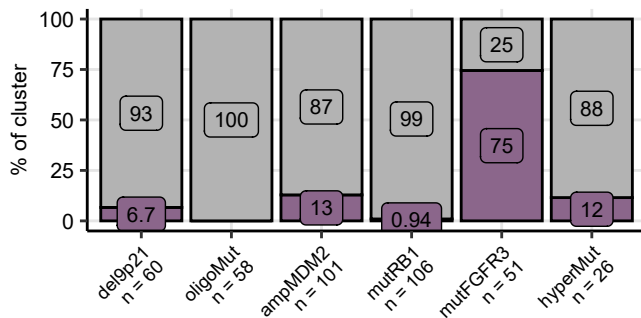
V = 0.79, p < 0.001

***FGFR3* mutation, MSK IMPACT**

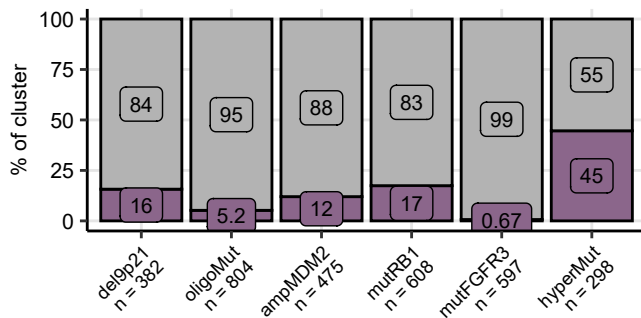
V = 0.76, p < 0.001

***FGFR3* mutation, TCGA BLCA**

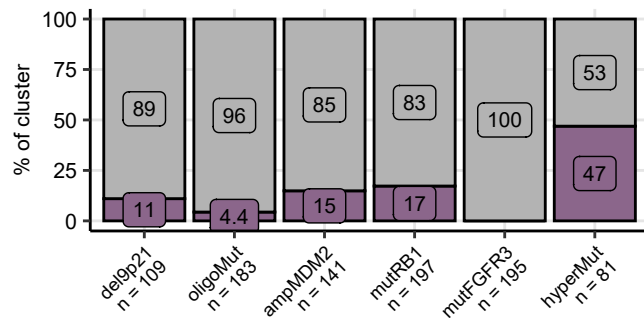
V = 0.65, p < 0.001

**B*****ERBB2* mutation, GENIE BLCA**

V = 0.36, p < 0.001

***ERBB2* mutation, MSK IMPACT**

V = 0.38, p < 0.001

***ERBB2* mutation, TCGA BLCA**

V = 0.32, p < 0.001

