

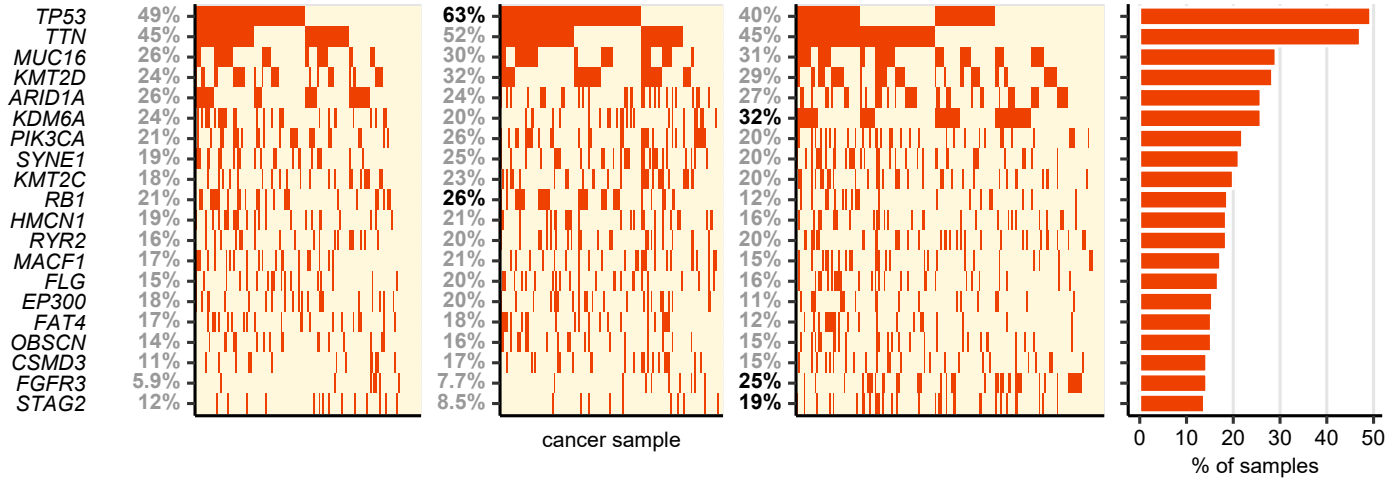
### TCGA BLCA, top most common mutations

#1: n = 119

#2: n = 117

#3: n = 169

cohort: n = 405



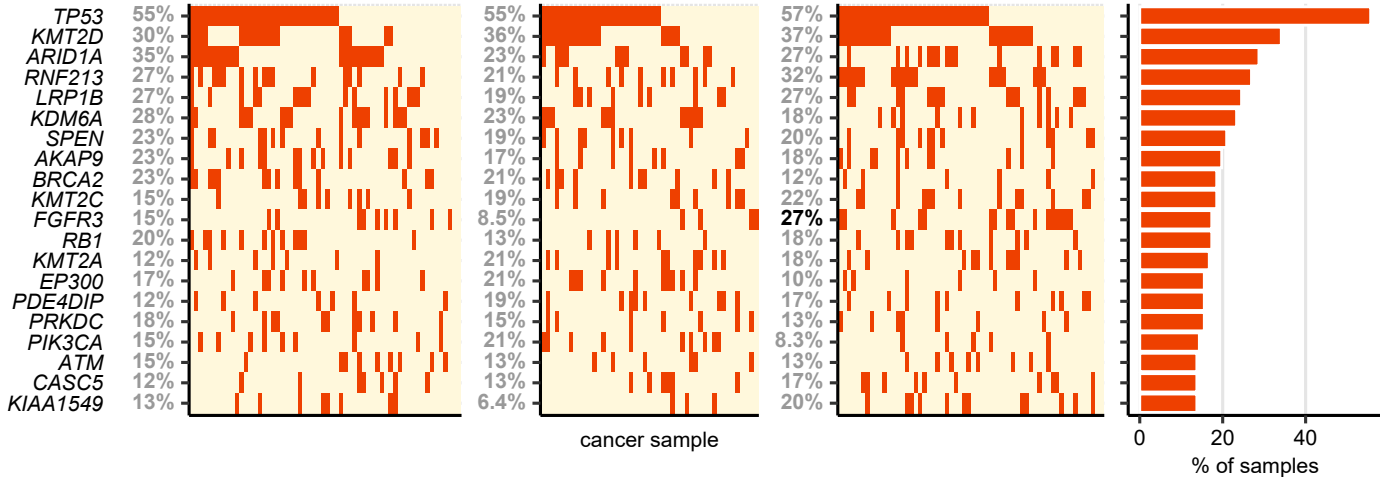
### BCAN, top most common mutations

#1: n = 60

#2: n = 47

#3: n = 60

cohort: n = 167



### IMvigor, top most common mutations

#1: n = 57

#2: n = 44

#3: n = 79

cohort: n = 180

