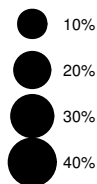


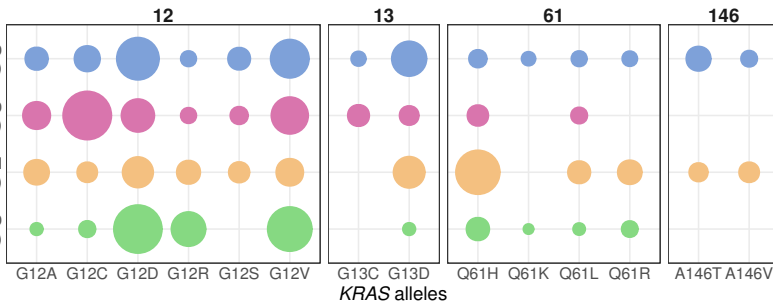
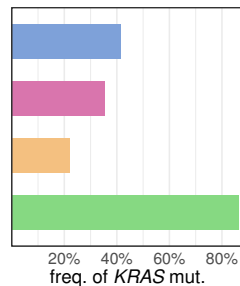
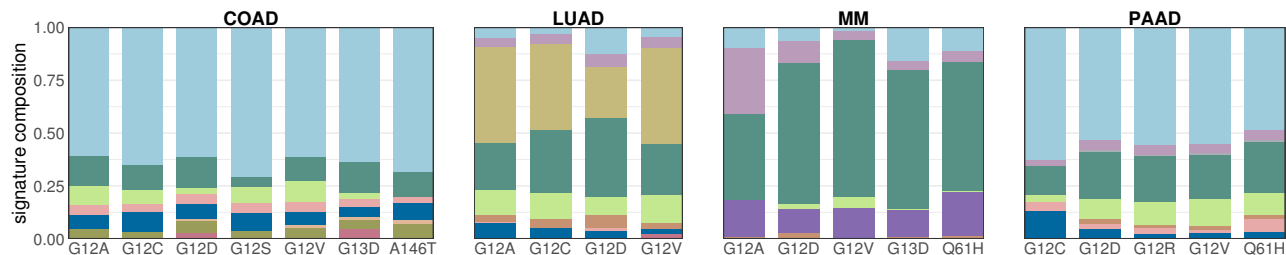
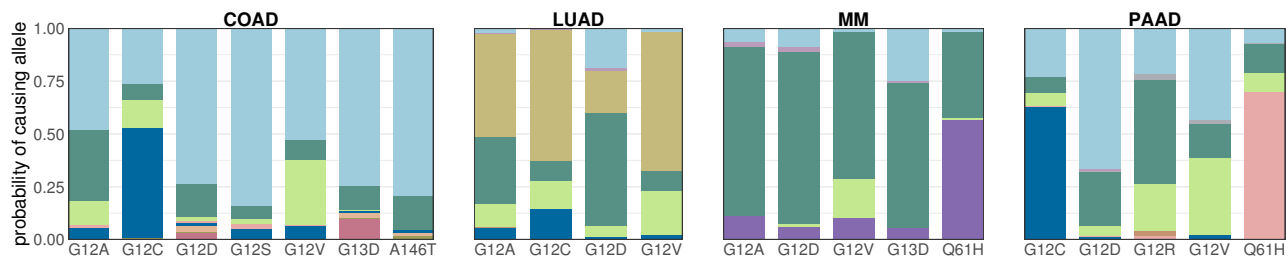
**a**percent of  
*KRAS* mutations

COAD  
(colon, n=4,145)

LUAD  
(lung, n=5,051)

MM  
(WBC, n=1,262)

PAAD  
(pancreas, n=2,314)

**b****c****d**

signature 1 2 3 4 5 8 9 13 17 18 30 N MSI