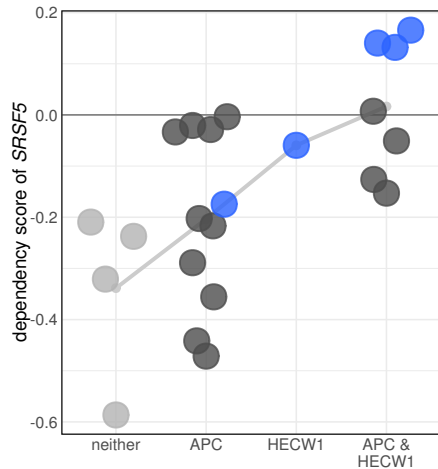
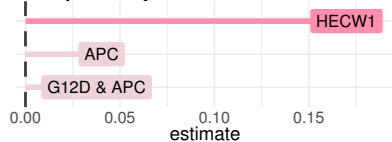


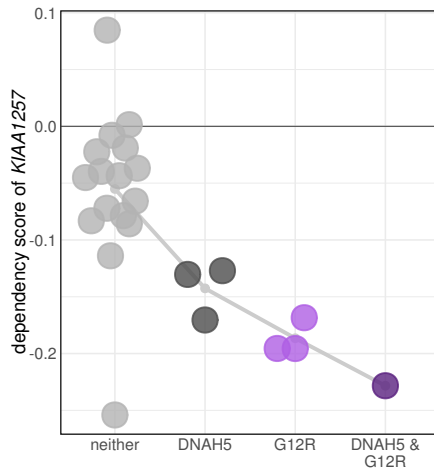
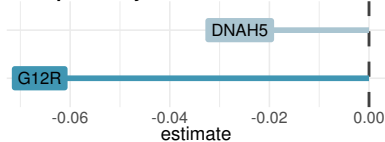
a COAD
estimated effect of mutation
on dependency on *SRSF5*



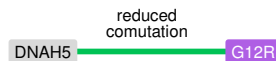
● G12D ● APC/HECW1 ● neither



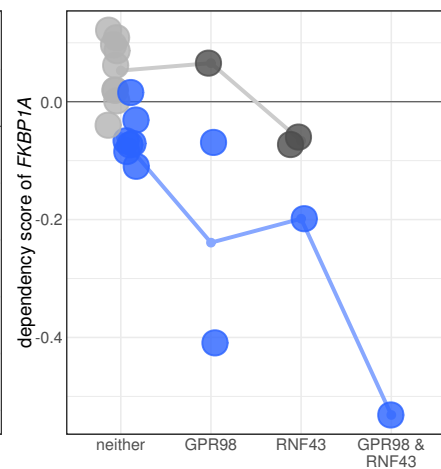
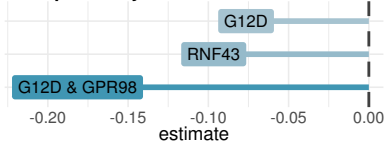
b PAAD
estimated effect of mutation
on dependency on *KIAA1257*



● neither ● DNAH5 ● G12R ● DNAH5 & G12R



c PAAD
estimated effect of mutation
on dependency on *FKBP1A*



● neither ● GPR98/RNF43 ● G12D

