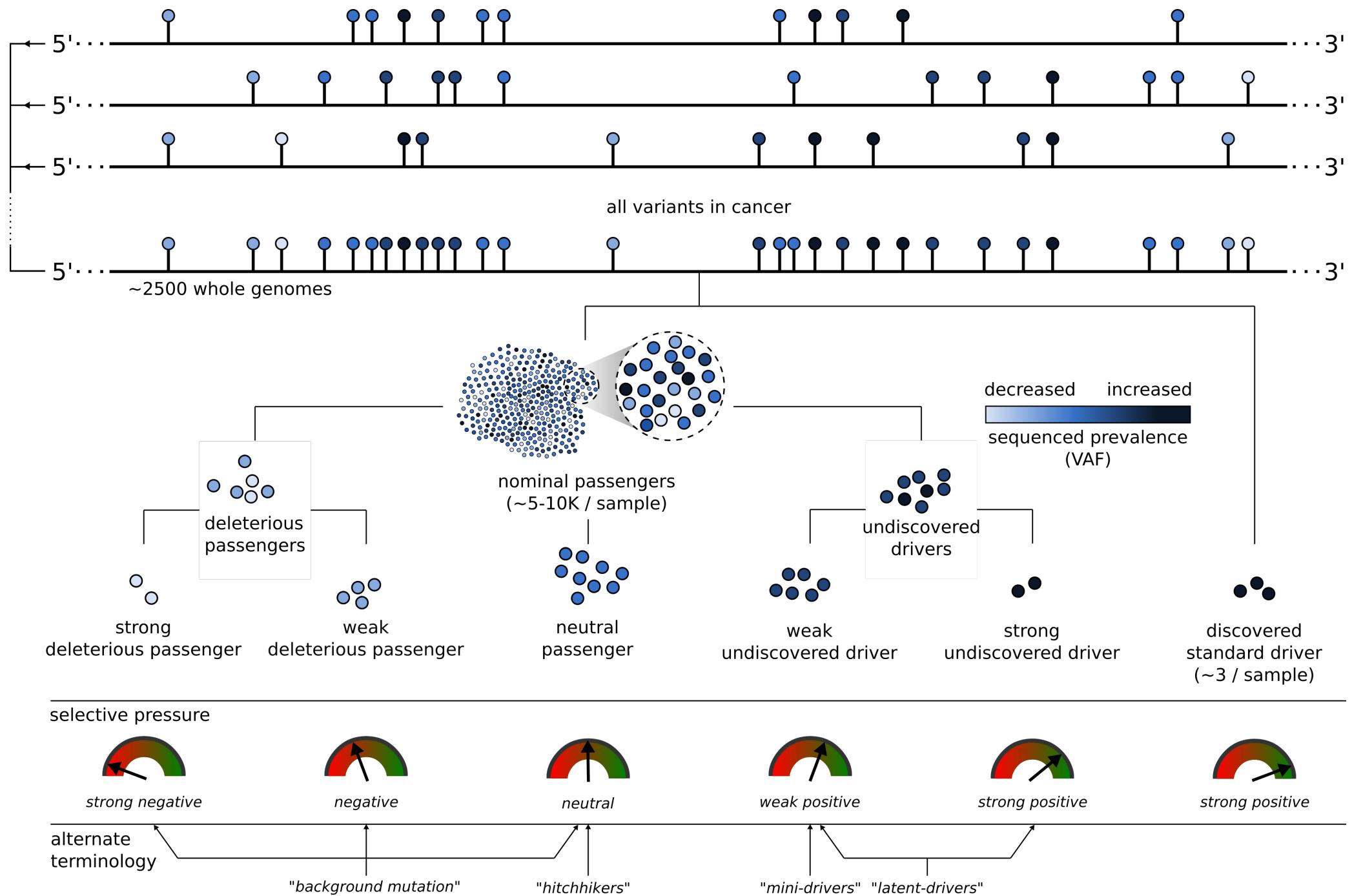
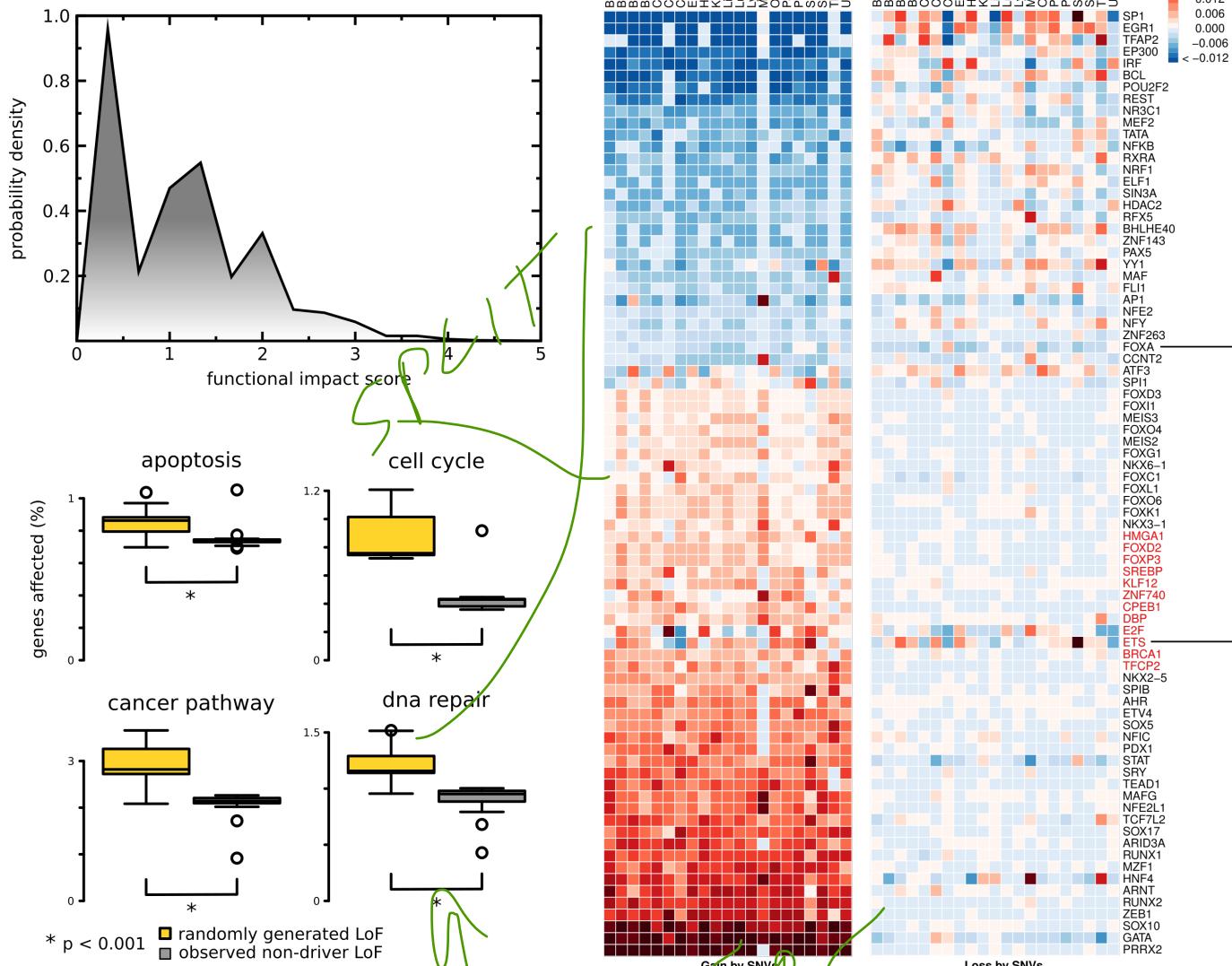
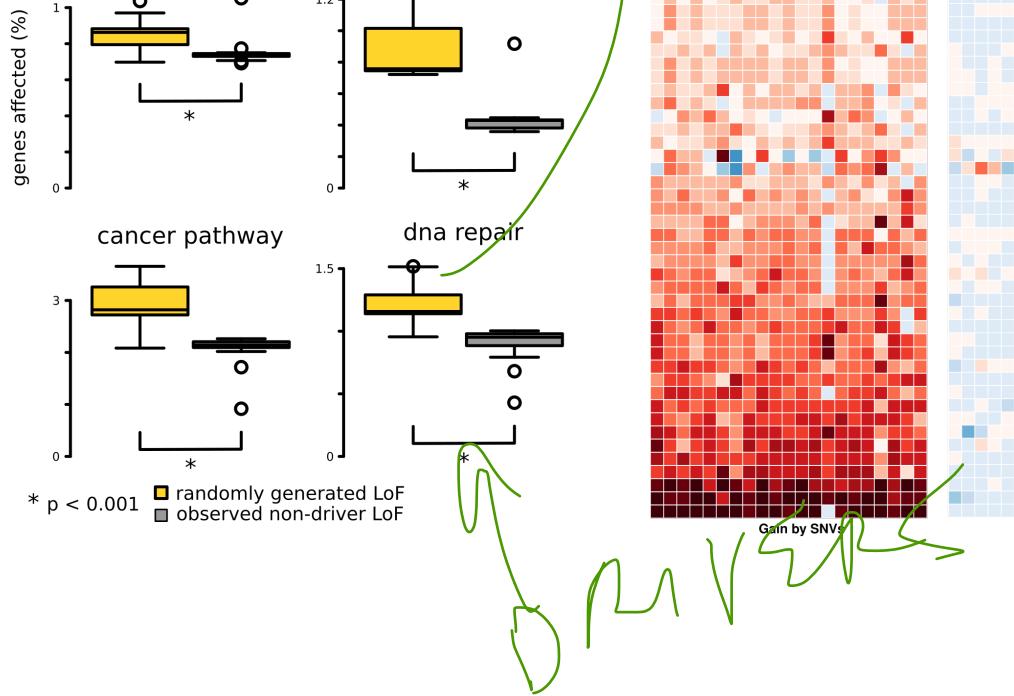
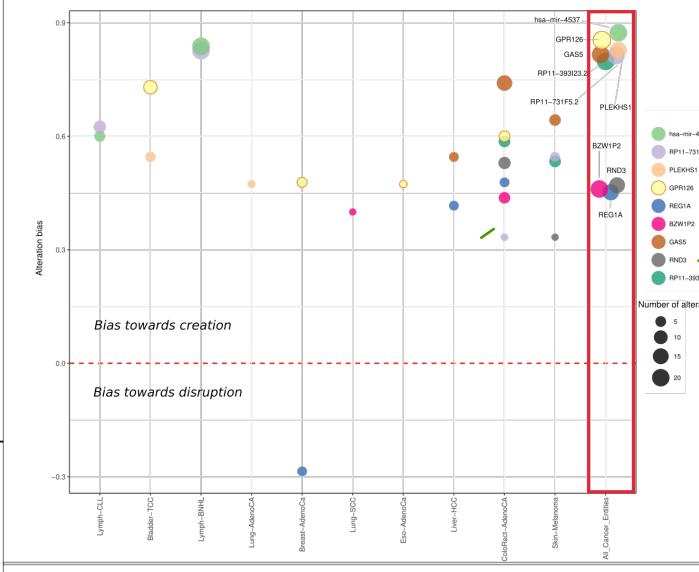
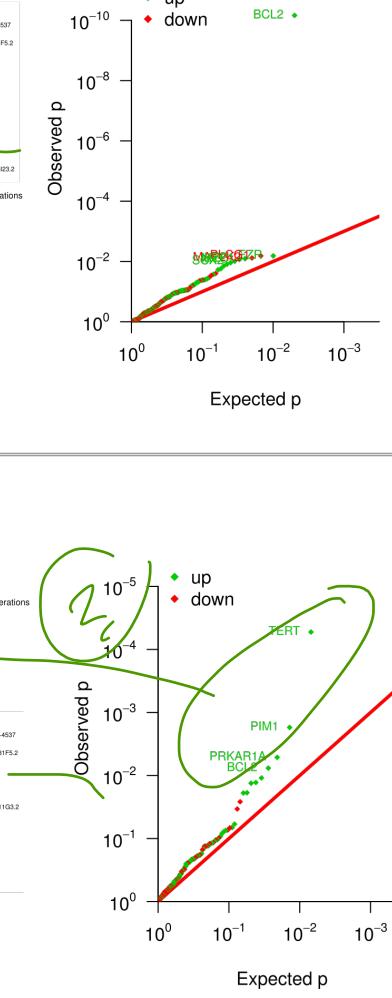
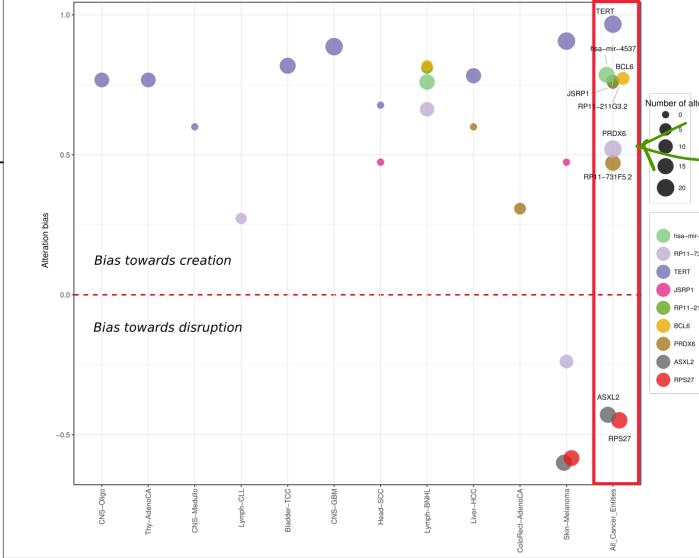
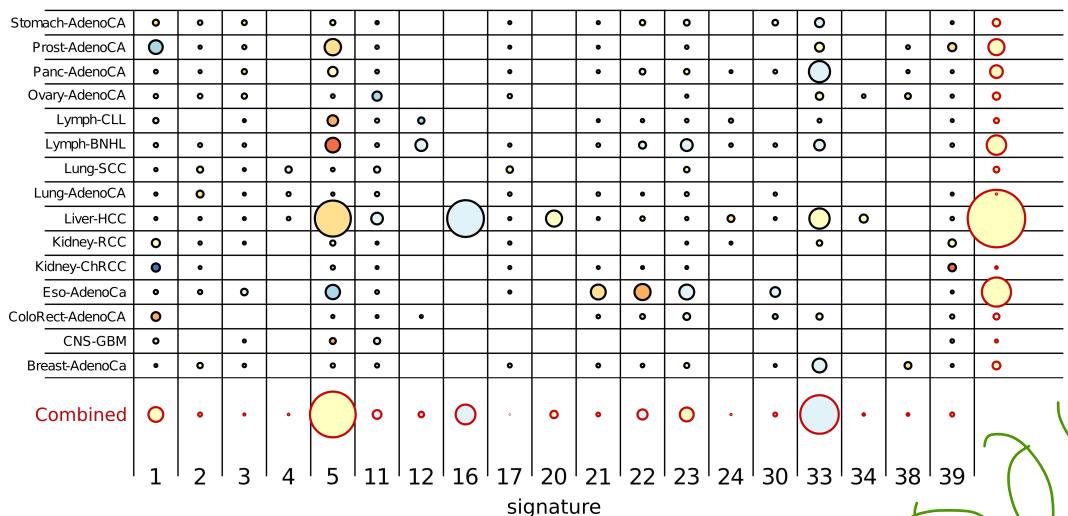
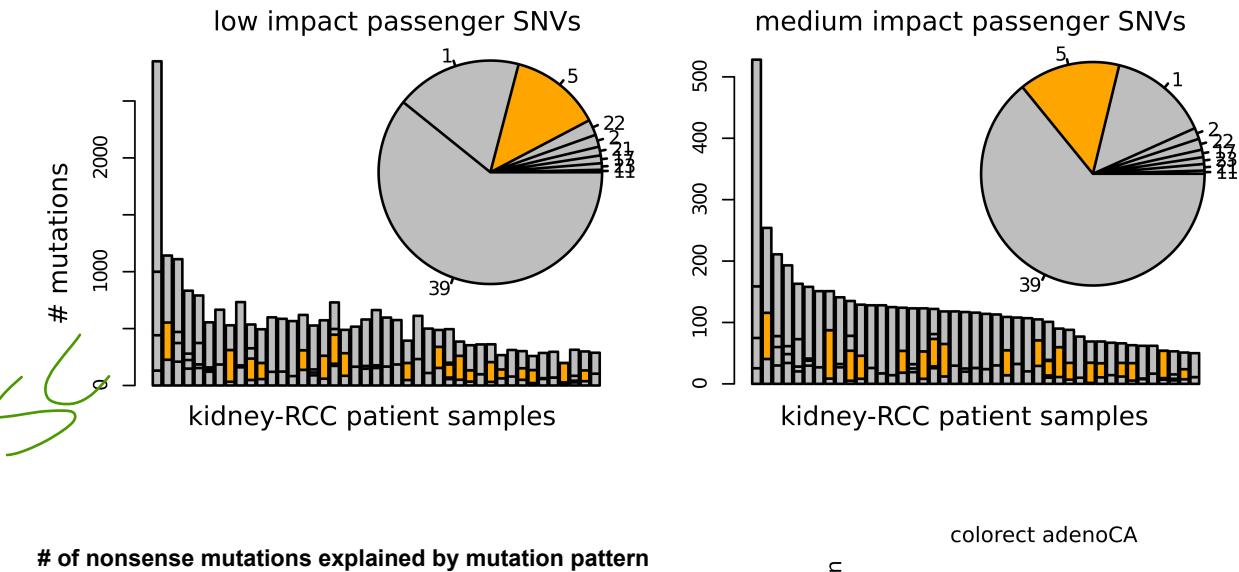
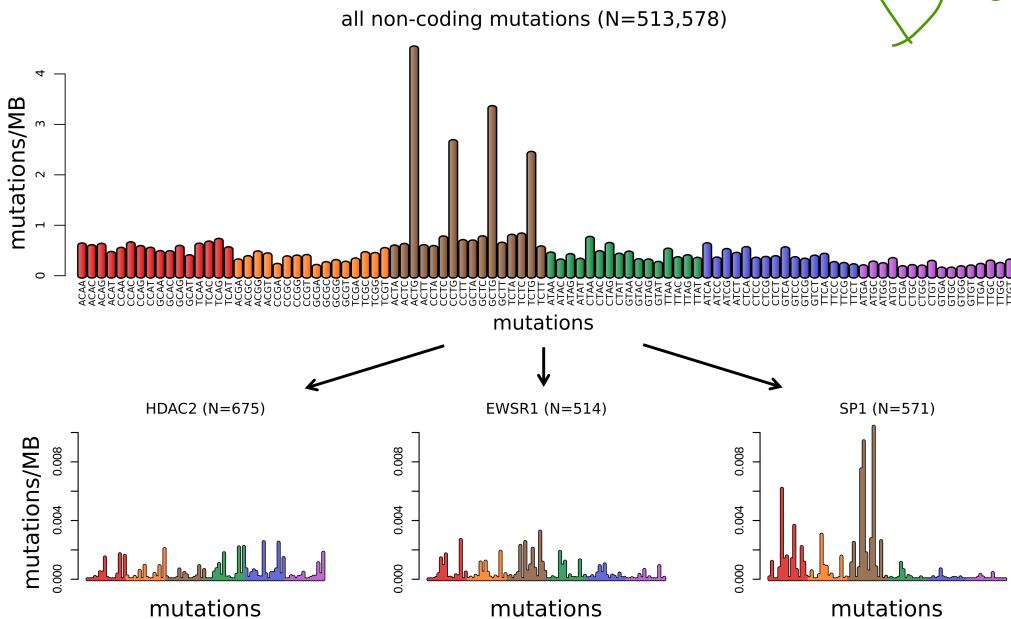


Figure Pack

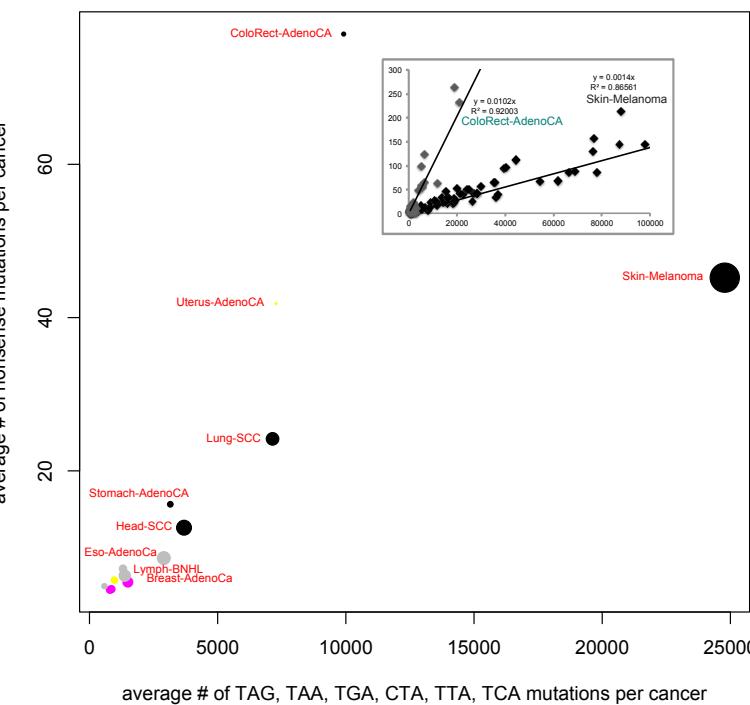
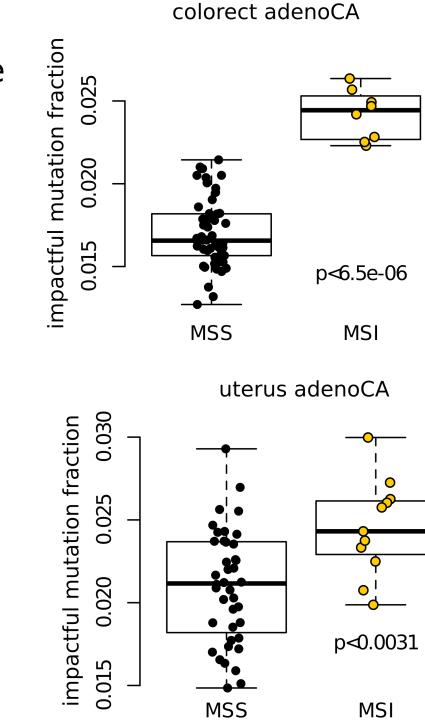
06/05/2017

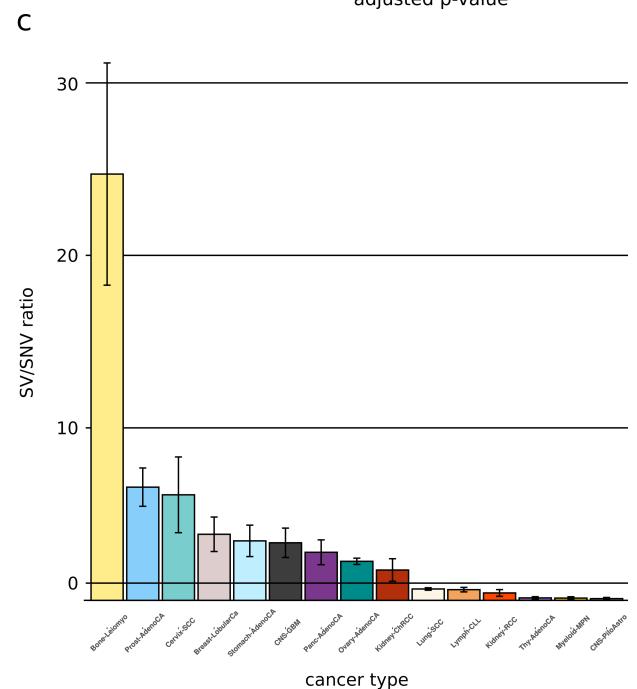
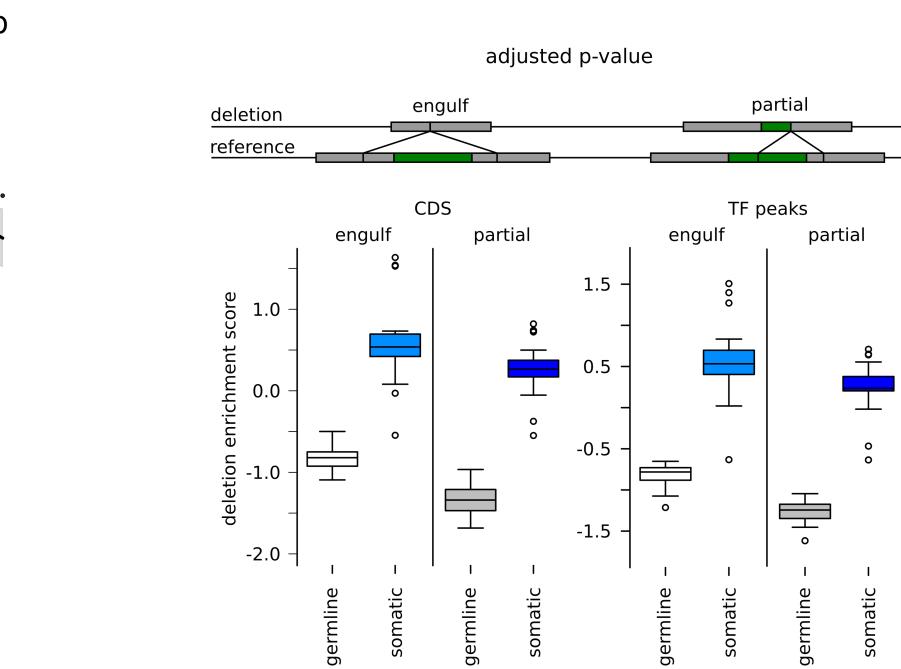
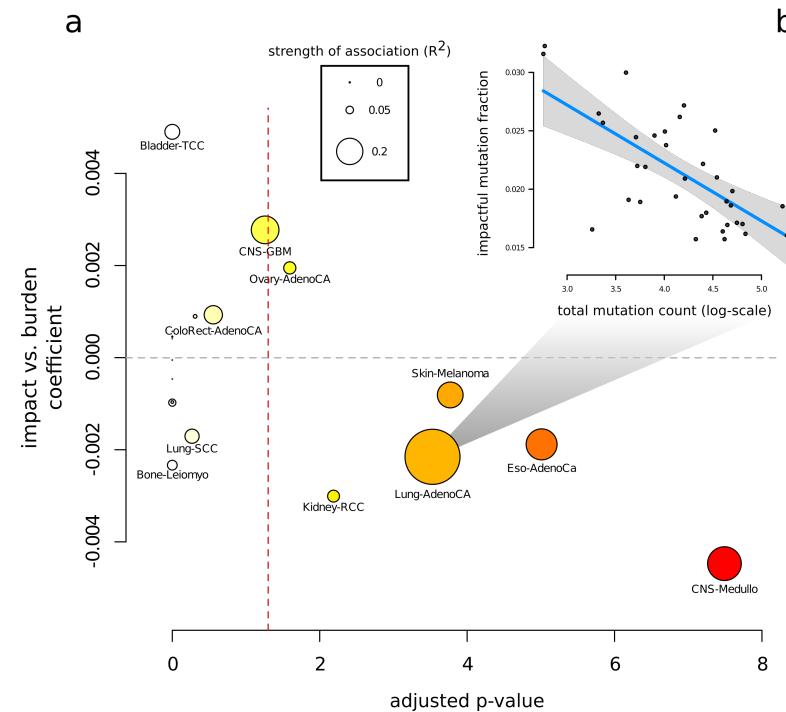


a**c****FOX family****ETS family**

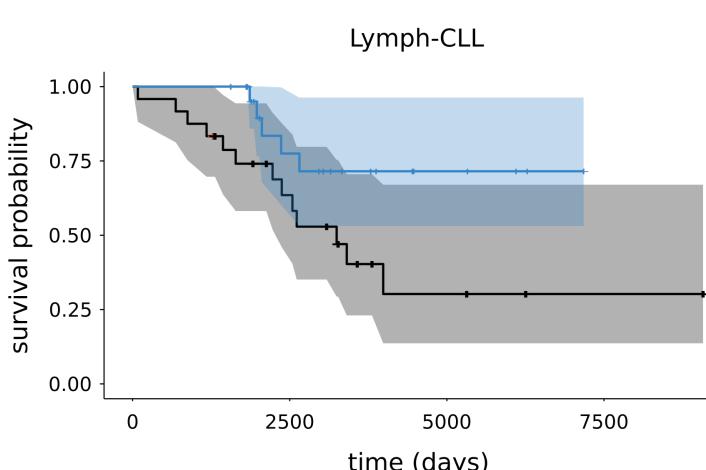
a**b****c**

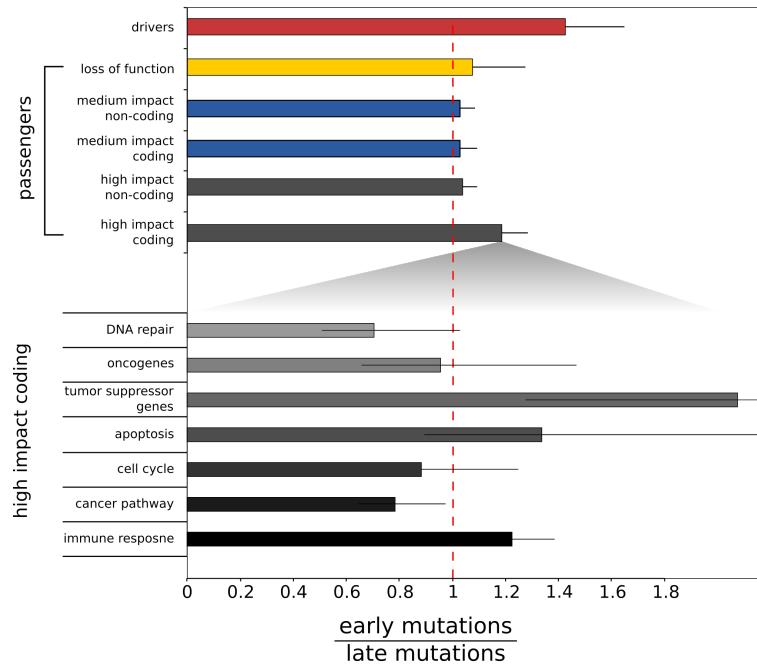
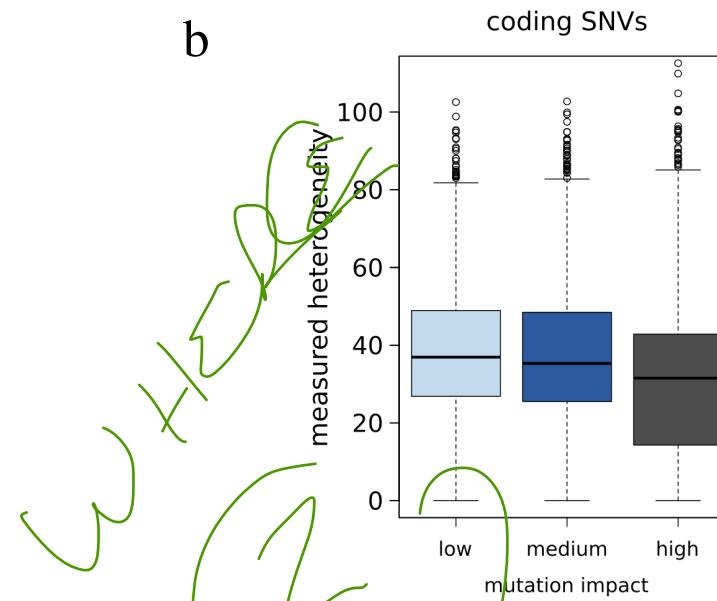
average # of nonsense mutations per cancer

**e**

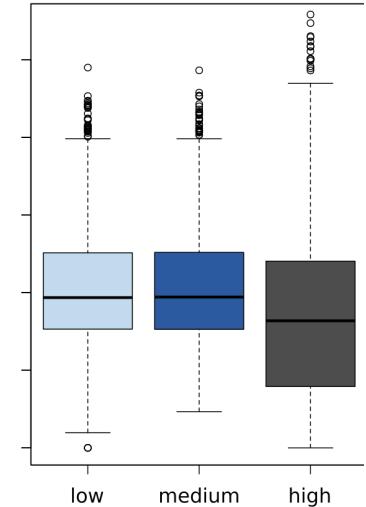
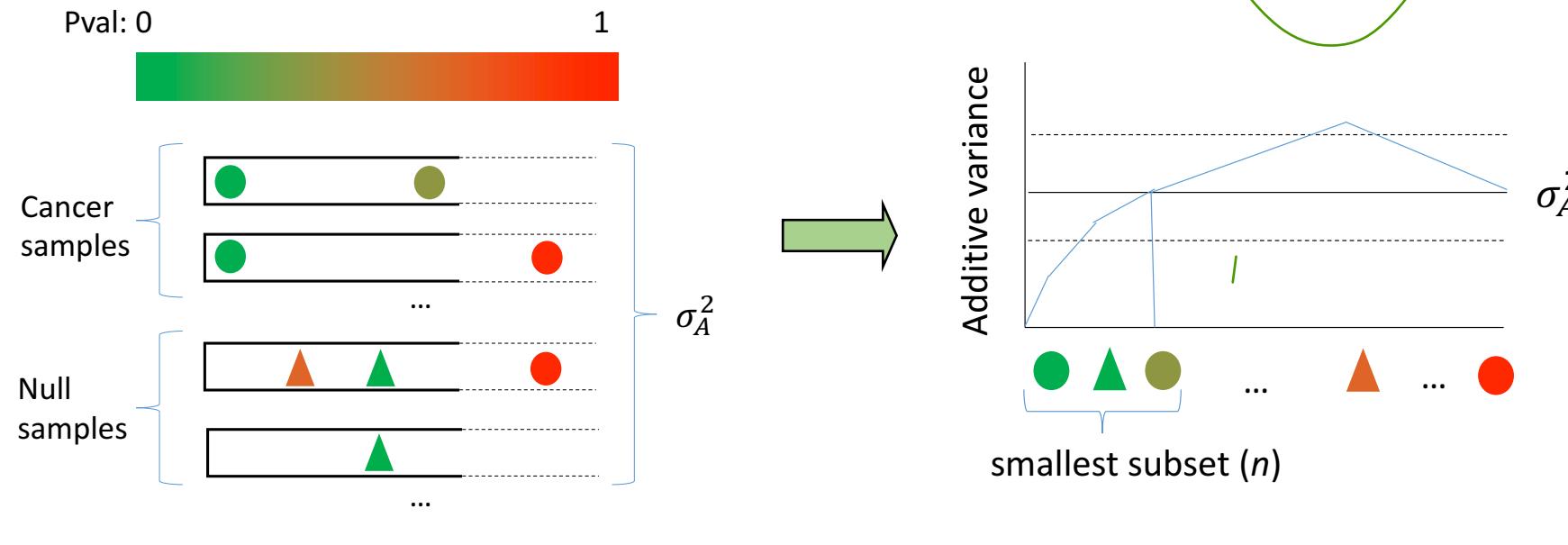


high impact burden low impact burden



a**b**

non-coding SNVs

**c**

Additive variance

smallest subset (n)