Altered in 356 (62.79%) of 567 samples. 1847 TMB 270 No. of samples TP53 48% **ATM** 8% **PRKDC** 8% **FANCM** 6% BRCA2 6% HFM1 6% POLE 5% **ATR** 4% BRCA1 4% MSH4 4% Splice_Site Translation_Start_Site Missense_Mutation In_Frame_Del

Nonsense_Mutation

Frame_Shift_Del

Frame Shift Ins

In_Frame_Ins

■ Multi_Hit