Altered in 38 (20.88%) of 182 samples. 2161 -TMB 9 No. of samples GPRASP1 3% APLP1 4% LING01 5% ALPK3 4% LZTS1 2% SPTBN5 3% 2% PCDHB14 LZTS3 4% RGL2 1% CYP4F12 3% Fustat Risk Missense_Mutation ■ Frame_Shift_Ins **Fustat** Risk Frame_Shift_Del Nonsense_Mutation Alive High In_Frame_Del Multi_Hit Dead Low