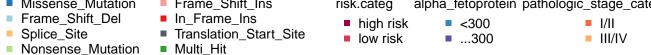
Altered in 264 (84.6153846153846%) of 312 samples. 1293 97 31% 27% 12% CTNNB1 ALB 10% 8% 7% 5% APOB ARID1A AXIN1 ARID2 BAP1 5% 5% 4% PS6KA3 KEAP1 4% 4% 4% 3% IDH1 CDKN1A CREB3L3 GNAS WHSC1 KRAS 1% 1% HST1H1E NRAS 1% **NUP113** 0% Missense Mutation Frame Shift Ins risk.categ alpha\_fetoprotein pathologic\_stage\_categ



In Frame Del