Altered in 68 (100%) of 68 samples. 162 TMB 67 No. of samples NCOR2 99% PIK3CA 40% APOB 28% BRCA2 26% KMT2C 22% TP53 22% FAT3 21% PKHD1 21% BIRC6 19% FAT1 19% CIC 18% KMT2D 16% LRP1B 16% PARP4 16% ASXL1 15% MAP3K1 15% NOTCH2 15% **POLE** 15% 15% RELN **APC** 13% **ATM** 13% EP300 13% FLT4 13% IGF2R 13% RANBP2 13% Tumor.location PAM50 PT Tumor.location PAM50 PT Missense Mutation Frame Shift Ins In Frame Ins Splice Site REC Basal LumB Nonsense\_Mutation Translation\_Start\_Site Her2 NA Frame\_Shift\_Del ■ Multi Hit LumA In\_Frame Del