Altered in 57 (100%) of 57 samples. 31 TMB 54 No. of samples 0 NCOR2 95% KMT2C 39% PKHD1 35% PALB2 23% FAT3 21% LRP1B 21% BRCA2 19% FOXC1 19% PIK3C2B 19% TET2 19% 18% **ATR** CIC 18% **GNAQ** 18% NCOR1 18% PARP4 18% **APOB** 16% NOTCH2 16% **RELN** 16% ASXL1 14% BCL9 14% BIRC6 14% NOTCH3 14% PIK3CA 14% RANBP2 14% **TERT** 14% Tumor.location PAM50 PT Missense Mutation In_Frame_Del Tumor.location PAM50 PT In Frame Ins Frame Shift Del LNM ■ Basal ■ LumB Frame_Shift_Ins Multi Hit Her2 NA Nonsense Mutation LumA