Altered in 22 (95.65%) of 23 samples. 187 0 0 KMT2C 39% AHNAK2 30% 26% 22% PDE4DIP ANKRD36C BAGE2 22% MST1L 22% ZIC2 22% 17% ADAMTSL4 17% 17% 17% EGFR FLT3LG FRG1 17% IRX5 17% MYT1L 17% PCMTD1 PRAMEF4 17% 17% RHPN2 17% RTP1 17% 17% SSPO ABCC5 13% AGXT AMMECR1 13% 13% ANAPC1 13% ASH1L 13% ATAD3B 13% 13% 13% C11orf80 COL6A3 CSPG4 13% CTAGE6 13% DMBT1 13% EVL 13% FBLN2 GPR32 13% 13% GSAP 13% HDAC2 13% LAMA5 13% MLLT6 13% MUC17 13% MUC4 13% 13% MYH13 NBPF12 13% NOTCH2NL 13% 13% 13% POU5F1B PRB2 PRRT2 13% 13% 13% SIGLEC11 SLFN13 WBP1 13% ZDHHC11 13% ZNF99 FBN2 13% C>TT>AT>C C>G T>G C>A Missense_Mutation In_Frame_Del Frame_Shift_Del In_Frame_Ins Nonsense_Mutation Splice_Site Frame_Shift_Ins ■ Multi_Hit