Altered in 153 (85.96%) of 178 samples. 13304 137 0 KRAS 77% TP53 63% SMAD4 22% CDKN2A 16% TTN 15% MUC16 7% RNF43 7% **GNAS** 6% ARID1A RYR1 5% In_Frame_Ins Nonsense_Mutation Missense_Mutation In_Frame_Del Frame_Shift_Del ■ Frame_Shift_Ins Splice_Site ■ Multi_Hit