Altered in 92 (93.88%) of 98 samples. 1143 -TMB 84 No. of samples TP53 86% TTN22% MUC16 13% 13% SYNE1 FAT3 12% SPTA1 12% CSMD3 10% DMD 10% DYNC2H1 10% PIK3CA 10% Cluster Cluster Missense_Mutation In_Frame_Ins ■ Nonsense_Mutation ■ Frame_Shift_Ins clust1 Frame_Shift_Del In_Frame_Del clust2 Splice_Site ■ Multi_Hit clust3