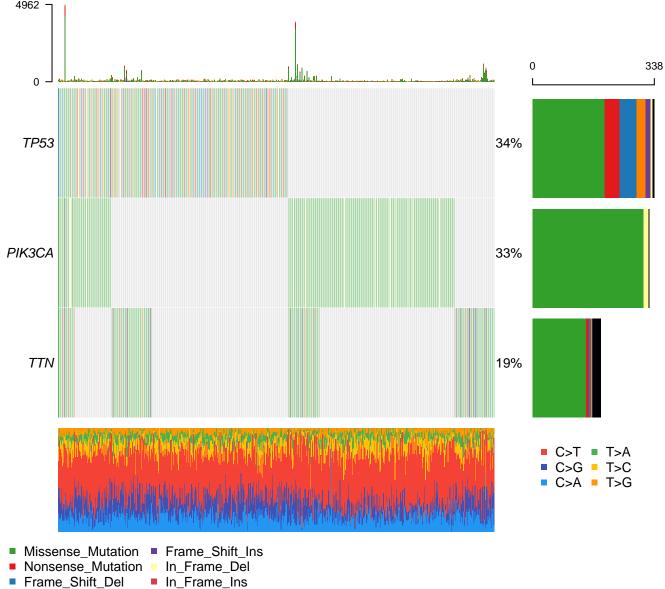
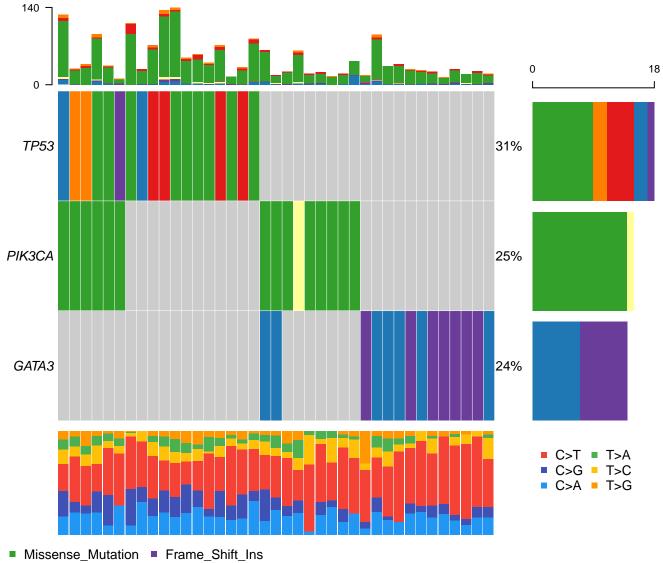
Altered in 643 (65.21%) of 986 samples.

Splice_Site

■ Multi_Hit



Altered in 39 (66.1%) of 59 samples.



- Splice_Site In_Frame_Del Nonsense_Mutation ■ Multi_Hit
- Frame_Shift_Del

Altered in 273 (69.11%) of 395 samples. 914 0 164 TP53 42% PIK3CA 30% TTN 17% ■ C>T ■ T>A C>G T>C C>A T>G Missense_Mutation Splice_Site



Altered in 258 (64.18%) of 402 samples. 4962 147 PIK3CA 37% TP53 29% TTN 24% ■ C>T ■ T>A C>G T>C C>A T>G



