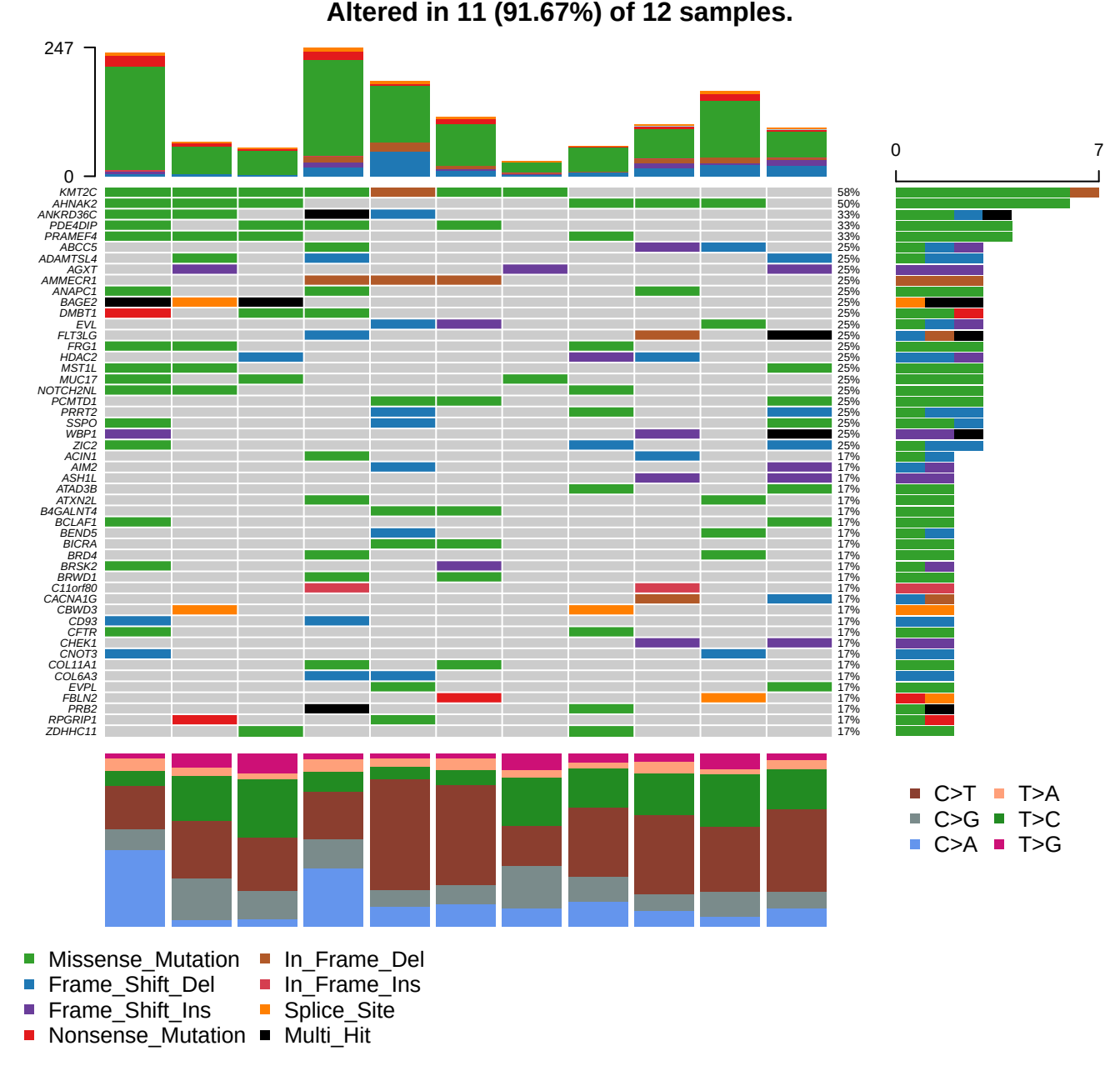
Altered in 11 (91.67%) of 12 samples.
247
0
0
7
KMT2C 58%
AHNAK2 50%
ANKRD36C 33%
PDE4DIP 33%
PRAMEF4 33%
ABCC5 25%
ADAMTSL4 25%
AGXT 25%
AMMECR1 25%
ANAPC1 25%
BAGE2 25%
DMBT1 25%
EVL 25%
FLT3LG 25%
FRG1 25%
HDAC2 25%
MST1L 25%
MUC17 25%
NOTCH2NL 25%
PCMTD1 25%
PRRT2 25%
SSPO 25%
WBP1 25%
ZIC2 25%
ACIN1 17%
AIM2 17%
ASH1L 17%
ATAD3B 17%
ATXN2L 17%
B4GALNT4 17%
BCLAF1 17%
BEND5 17%
BICRA 17%
BRD4 17%
BRSK2 17%
BRWD1 17%
C11orf80 17%
CACNA1G 17%
CBWD3 17%
CD93 17%
CFTR 17%
CHEK1 17%
CNOT3 17%
COL11A1 17%
COL6A3 17%
EVPL 17%
FBLN2 17%
PRB2 17%
RPGRIP1 17%
ZDHHC11 17%
C>T
C>G
C>A
T>A
T>C
T>G
Missense_Mutation
Frame_Shift_Del
Frame_Shift_Ins
Nonsense_Mutation
In_Frame_Del
In_Frame_Ins
Splice_Site
Multi_Hit