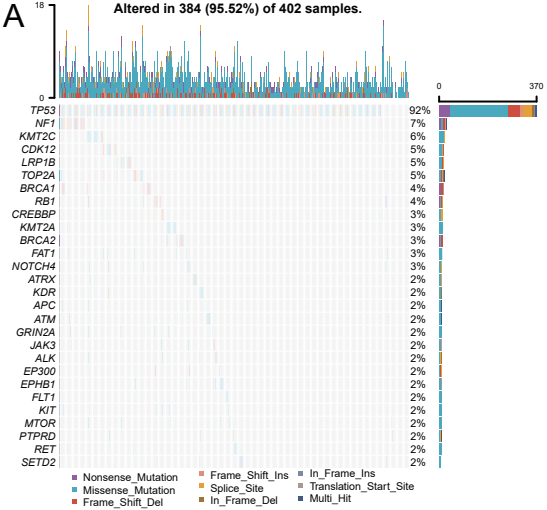
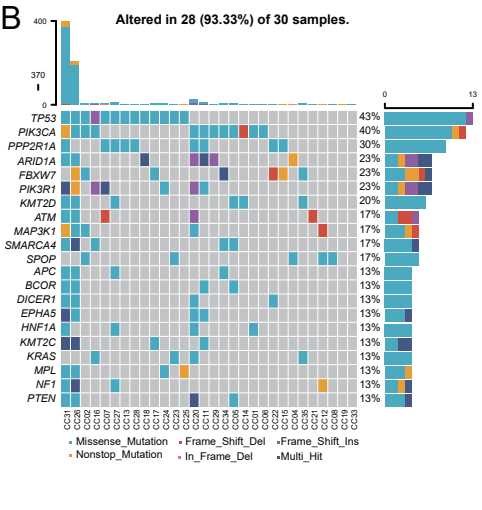

A
Altered in 384 (95.52%) of 402 samples.
18
0
0
370
TP53 92%
NF1 7%
KMT2C 6%
CDK12 5%
LRP1B 5%
TOP2A 5%
BRCA1 4%
RB1 4%
CREBBP 3%
KMT2A 3%
BRCA2 3%
FAT1 3%
NOTCH4 3%
ATRX 2%
KDR 2%
APC 2%
ATM 2%
GRIN2A 2%
JAK3 2%
ALK 2%
EP300 2%
EPHB1 2%
FLT1 2%
KIT 2%
MTOR 2%
PTPRD 2%
RET 2%
SETD2 2%
Nonsense_Mutation
Missense_Mutation
Frame_Shift_Del
Frame_Shift_Ins
Splice_Site
In_Frame_Del
In_Frame_Ins
Translation_Start_Site
Multi_Hit


B
Altered in 28 (93.33%) of 30 samples.
400
370
0
0
13
TP53 43%
PIK3CA 40%
PPP2R1A 30%
ARID1A 23%
FBXW7 23%
PIK3R1 23%
KMT2D 20%
ATM 17%
MAP3K1 17%
SMARCA4 17%
SPOP 17%
APC 13%
BCOR 13%
DICER1 13%
EPHA5 13%
HNF1A 13%
KMT2C 13%
KRAS 13%
MPL 13%
NF1 13%
PTEN 13%
CC13 CC26 CC02 CC16 CC17 CC27 CC13 CC18 CC18 CC17 CC24 CC25 CC25 CC20 CC17 CC29 CC14 CC16 CC14 CC01 CC06 CC07 CC15 CC04 CC21 CC21 CC12 CC08 CC28 CC03
Missense_Mutation
Nonstop_Mutation
Frame_Shift_Del
In_Frame_Del
Frame_Shift_Ins
Multi_Hit