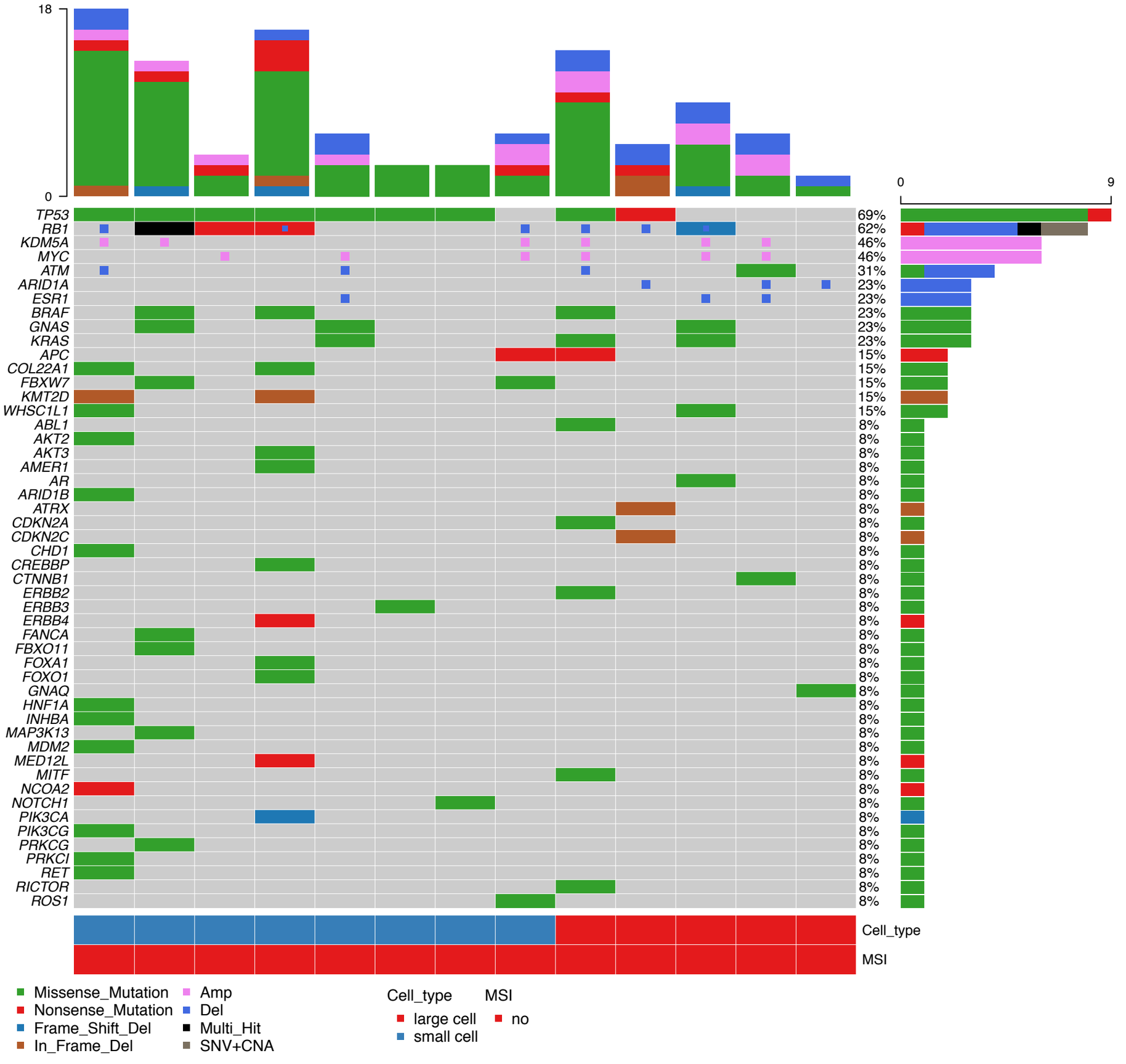
18
0
0
9
TP53 69%
RB1 62%
KDM5A 46%
MYC 46%
ATM 31%
ARID1A 23%
ESR1 23%
BRAF 23%
GNAS 23%
KRAS 23%
APC 15%
COL22A1 15%
FBXW7 15%
KMT2D 15%
WHSC1L1 15%
ABL1 8%
AKT2 8%
AKT3 8%
AMER1 8%
AR 8%
ARID1B 8%
ATRX 8%
CDKN2A 8%
CDKN2C 8%
CHD1 8%
CREBBP 8%
CTNNB1 8%
ERBB2 8%
ERBB3 8%
ERBB4 8%
FANCA 8%
FBXO11 8%
FOXA1 8%
FOXO1 8%
GNAQ 8%
HNF1A 8%
INHBA 8%
MAP3K13 8%
MDM2 8%
MED12L 8%
MITF 8%
NCOA2 8%
NOTCH1 8%
PIK3CA 8%
PIK3CG 8%
PRKCG 8%
PRKCI 8%
RET 8%
RICTOR 8%
ROS1 8%
Cell_type
MSI
Missense_Mutation
Nonsense_Mutation
Frame_Shift_Del
In_Frame_Del
Amp
Del
Multi_Hit
SNV+CNA
Cell_type
large cell
small cell
MSI
no