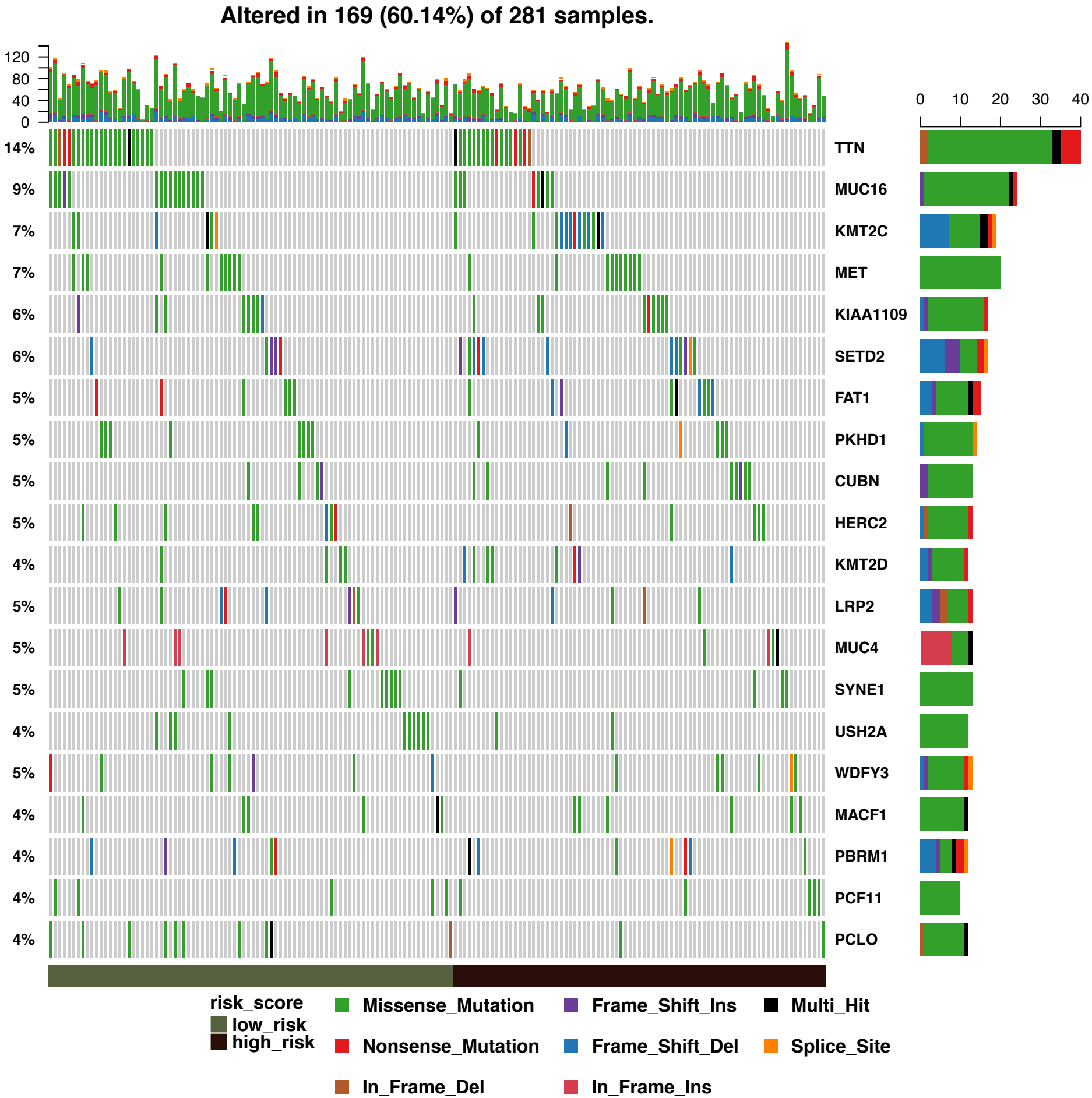
Altered in 169 (60.14%) of 281 samples.
120
80
40
0
0
10
20
30
40
14%
TTN
9%
MUC16
7%
KMT2C
7%
MET
6%
KIAA1109
6%
SETD2
5%
FAT1
5%
PKHD1
5%
CUBN
5%
HERC2
4%
KMT2D
5%
LRP2
5%
MUC4
5%
SYNE1
4%
USH2A
5%
WDFY3
4%
MACF1
4%
PBRM1
4%
PCF11
4%
PCLO
risk_score
low_risk
high_risk
Missense_Mutation
Nonsense_Mutation
In_Frame_Del
Frame_Shift_Ins
Frame_Shift_Del
In_Frame_Ins
Multi_Hit
Splice_Site