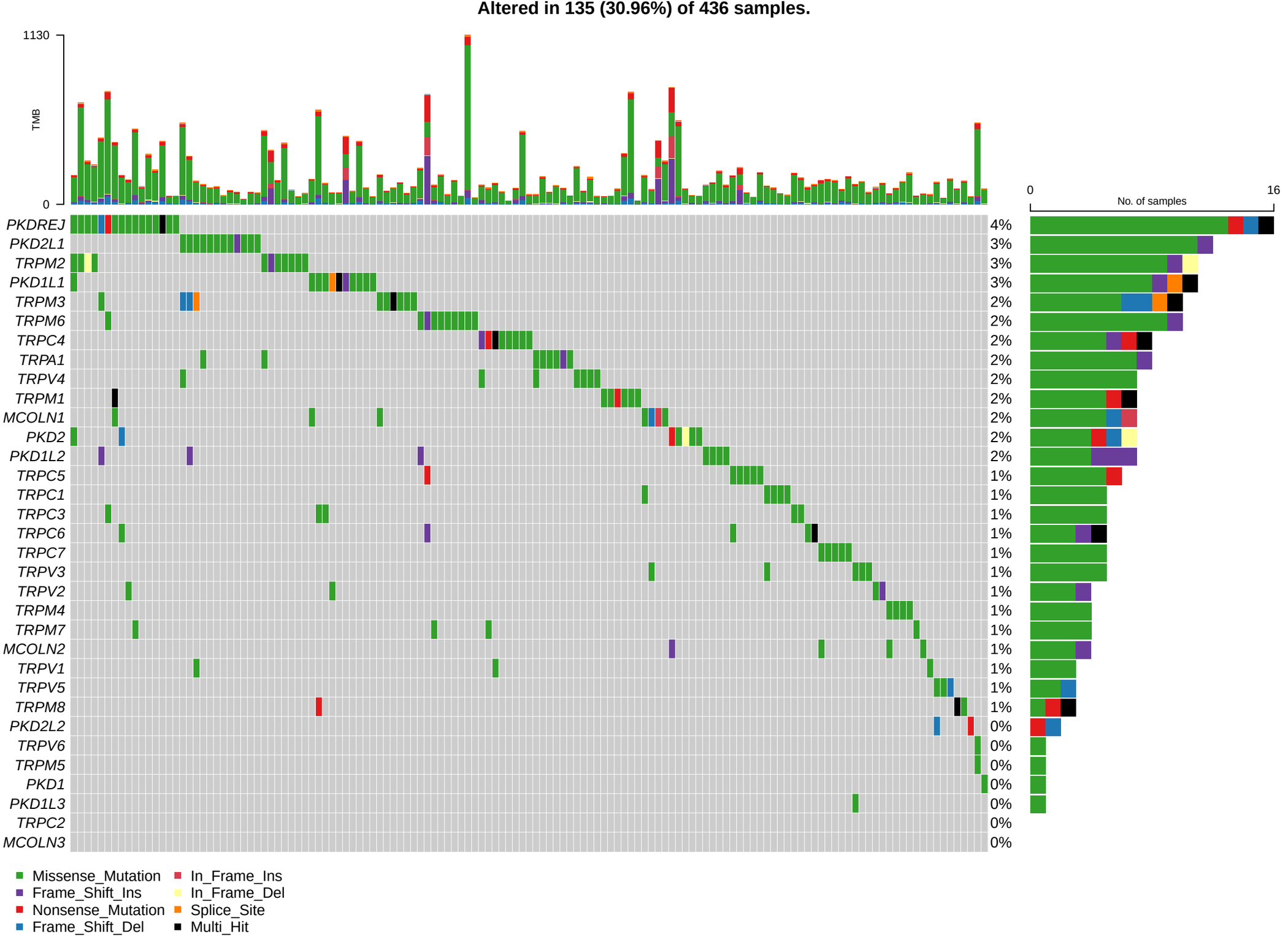
Altered in 135 (30.96%) of 436 samples.
1130
TMB
0
0
16
No. of samples
PKDREJ 4%
PKD2L1 3%
TRPM2 3%
PKD1L1 3%
TRPM3 2%
TRPM6 2%
TRPC4 2%
TRPA1 2%
TRPV4 2%
TRPM1 2%
MCOLN1 2%
PKD2 2%
PKD1L2 2%
TRPC5 1%
TRPC1 1%
TRPC3 1%
TRPC6 1%
TRPC7 1%
TRPV3 1%
TRPV2 1%
TRPM4 1%
TRPM7 1%
MCOLN2 1%
TRPV1 1%
TRPV5 1%
TRPM8 1%
PKD2L2 0%
TRPV6 0%
TRPM5 0%
PKD1 0%
PKD1L3 0%
TRPC2 0%
MCOLN3 0%
Missense_Mutation
Frame_Shift_Ins
Nonsense_Mutation
Frame_Shift_Del
In_Frame_Ins
In_Frame_Del
Splice_Site
Multi_Hit