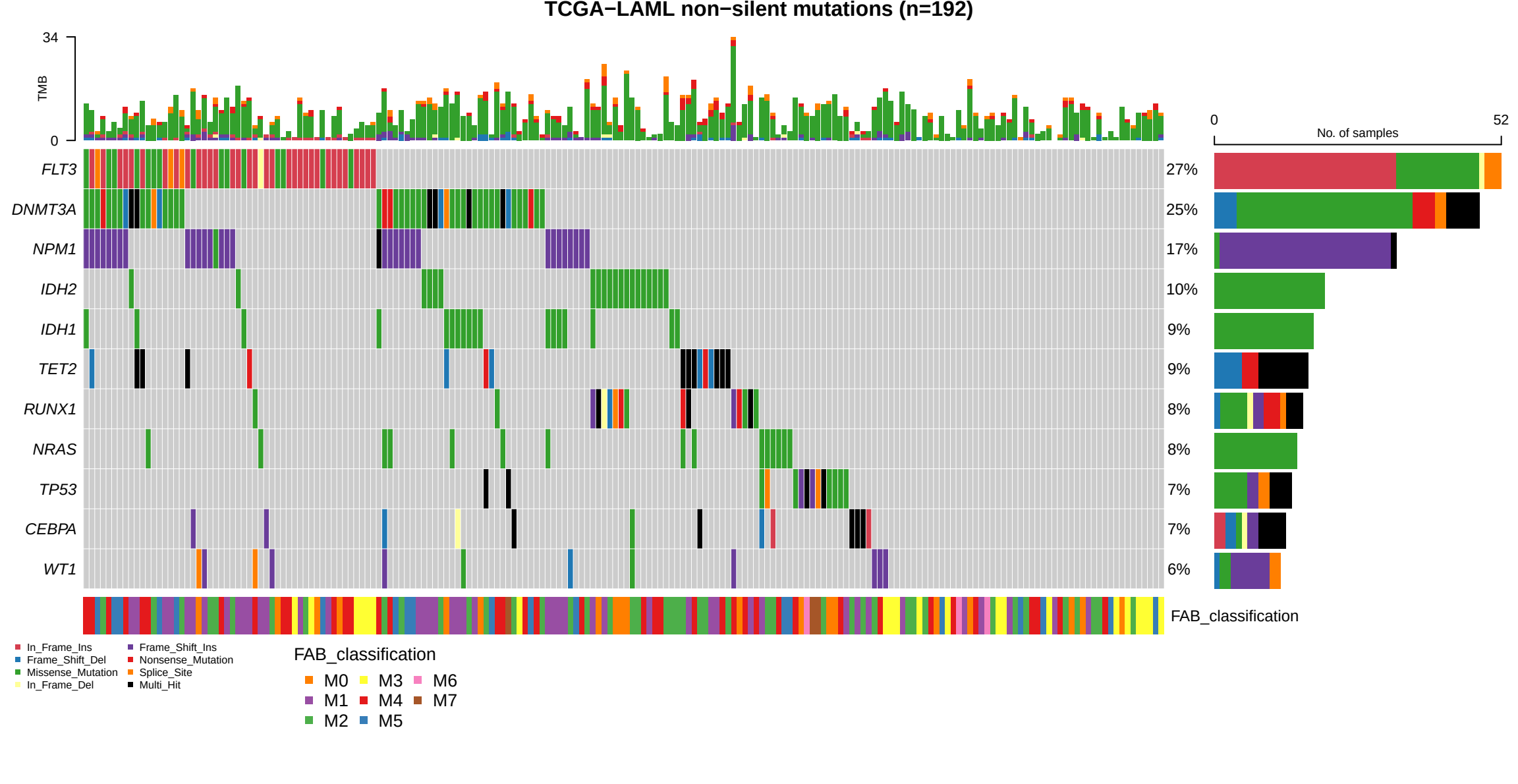

TCGA−LAML non−silent mutations (n=192)
34
TMB
0
0
52
No. of samples
FLT3
DNMT3A
NPM1
IDH2
IDH1
TET2
RUNX1
NRAS
TP53
CEBPA
WT1
27%
25%
17%
10%
9%
9%
8%
8%
7%
7%
6%
FAB_classification
In_Frame_Ins
Frame_Shift_Ins
Frame_Shift_Del
Nonsense_Mutation
Missense_Mutation
Splice_Site
In_Frame_Del
Multi_Hit
FAB_classification
M0
M3
M6
M1
M4
M7
M2
M5