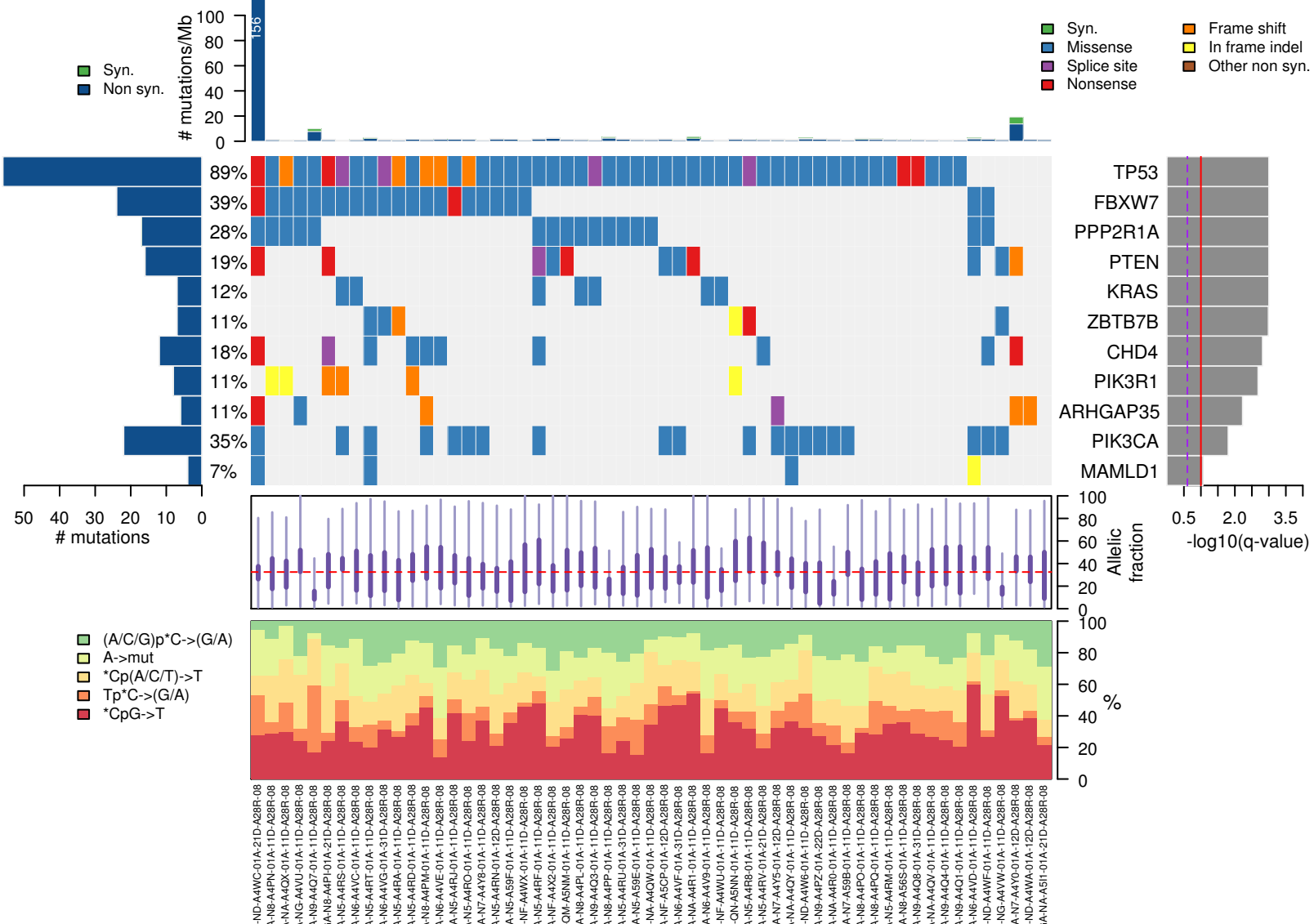
Syn.
Non syn.
mutations/Mb
156
Syn.
Missense
Splice site
Nonsense
Frame shift
In frame indel
Other non syn.
89%
39%
28%
19%
12%
11%
18%
11%
11%
35%
7%
TP53
FBXW7
PPP2R1A
PTEN
KRAS
ZBTB7B
CHD4
PIK3R1
ARHGAP35
PIK3CA
MAMLD1
mutations
-log10(q-value)
Allelic fraction
(A/C/G)p*C->(G/A)
A->mut
*Cp(A/C/T)->T
Tp*C->(G/A)
*CpG->T
%