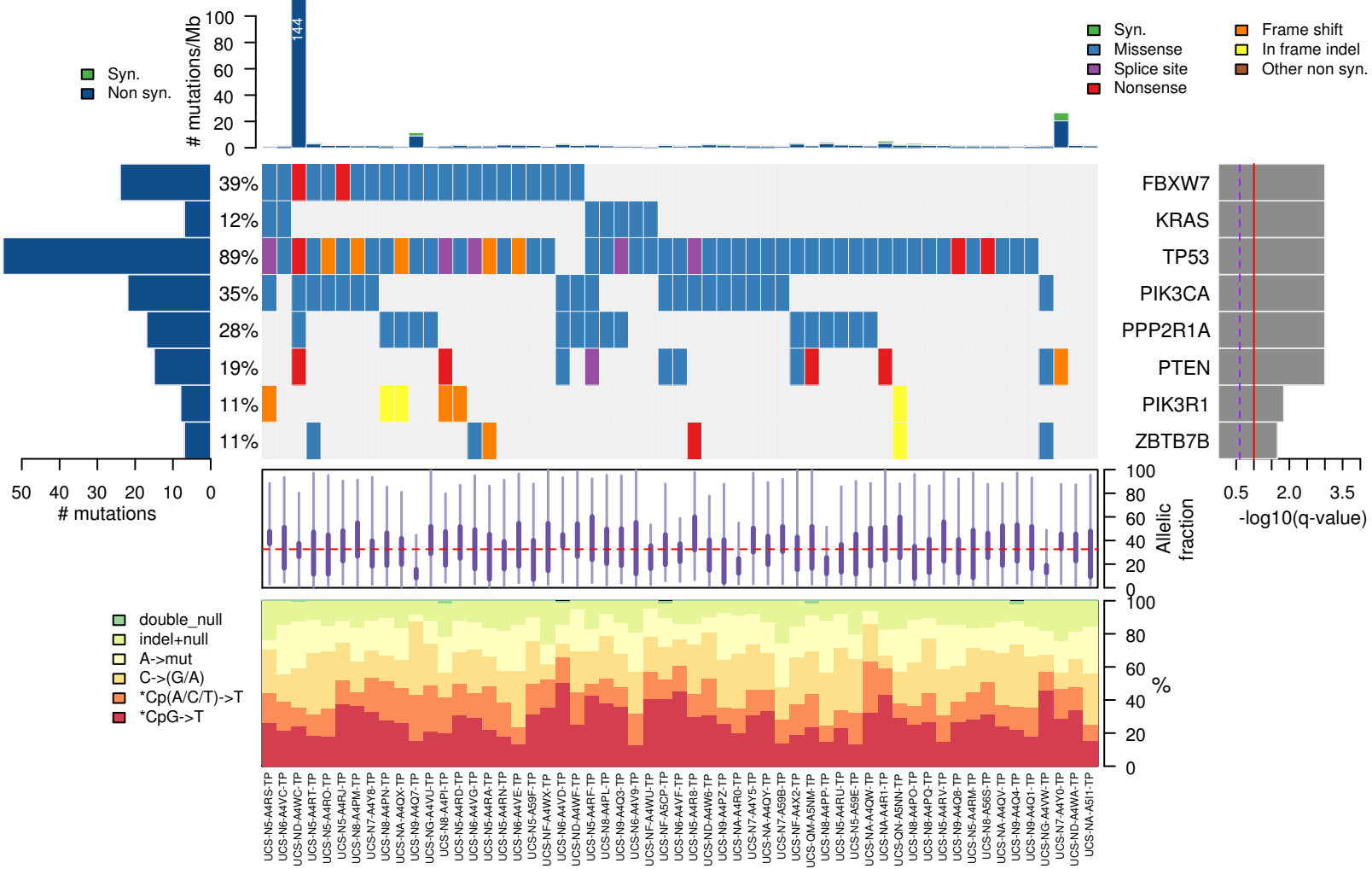
Syn.
Non syn.
mutations/Mb
144
Syn.
Missense
Splice site
Nonsense
Frame shift
In frame indel
Other non syn.
39%
12%
89%
35%
28%
19%
11%
11%
FBXW7
KRAS
TP53
PIK3CA
PPP2R1A
PTEN
PIK3R1
ZBTB7B
mutations
-log10(q-value)
Allelic fraction
double_null
indel+null
A->mut
C->(G/A)
*Cp(A/C/T)->T
*CpG->T
%